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Title:
Perfect score:
Sequence:
                                                                                                                                                                                                                          Result
No.
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Maximum Match 100%
Listing first 45 summaries
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Minimum DB seq length: 0
Maximum DB seq length: 2000000000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            OM protein - protein search, using sw model
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                                                                                                                                                                                                                                                                                                            Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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4::

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110::

111::

113::

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118::

118::
                                                                                                                                                                                                                      Match
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99.6
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1: /SIDS8/gcgdat
2: /SIDS8/gcgdat
                                                                                                                                                                                                                                                                                                                                                                             1: /SIDSB/gcgdata/geneseq/geneseqp/AA1980.DAT:*
2: /SIDSB/gcgdata/geneseq/geneseqp/AA1981.DAT:*
3: /SIDSB/gcgdata/geneseq/geneseqp/AA1981.DAT:*
4: /SIDSB/gcgdata/geneseq/geneseqp/AA1982.DAT:*
4: /SIDSB/gcgdata/geneseq/geneseqp/AA1983.DAT:*
5: /SIDSB/gcgdata/geneseq/geneseqp/AA1984.DAT:*
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7: /SIDSB/gcgdata/geneseq/geneseqp/AA1987.DAT:*
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10: /SIDSB/gcgdata/geneseq/geneseqp/AA1990.DAT:*
11: /SIDSB/gcgdata/geneseq/geneseqp/AA1992.DAT:*
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13: /SIDSB/gcgdata/geneseq/geneseqp/AA1993.DAT:*
14: /SIDSB/gcgdata/geneseq/geneseqp/AA1995.DAT:*
15: /SIDSB/gcgdata/geneseq/geneseqp/AA1995.DAT:*
16: /SIDSB/gcgdata/geneseq/geneseqp/AA1997.DAT:*
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19: /SIDSB/gcgdata/geneseq/geneseqp/AA1999.DAT:*
20: /SIDSB/gcgdata/geneseq/geneseqp/AA1999.DAT:*
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Gapop 10.0 , Gapext 0.5
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Copyright (c) 1993 - 2000 Comp
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AAY28408
AAY92236
AAY92257
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AAB35260
AAB352664
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AAB352664
AAB35263
AAB35263
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                              Human IL-1 recepto
Human IL-1 homolog
Human TANGO-93 pro
Human PRO4342. Ho
Human IL-IL1. Hom
Interleukin-IL1 re
Protein encoded by
A human interleuki
                                                                                                                                                                                                                      Description
                                                                                                                                                                                  Human interleukin
Interleukin-IL1
Interleukin-IL1
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319.5 38.8	.5 38.	.5 38.	.5 38.	.5 38.	.5 38.	.5 38.	.5 38.	.5 38.	.5 38	.5 38.	.5 38.	.5 38.	.5 38.	.5 38.	.5 38.	ഗ	98 48.	51.	53.	53.	63.	734 89.2	89.	89.	89.	89.	89.	89.	89.	97.	97.	9	816 99.1
165	159	159	159	159	159	159	159	159	159	159	153	153	153	152	152	159	98	94	80	80	104	156	156	156	156	155	155	155	155	155	155	155	155
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AAP93616	67	AAB14829	AAW83008	39	AAY28287	327	327	AAW37787	7	ιп	AAB51302	AAW61149	AAW22894	AAR35486	AAR27495	AAR73642	AAY97067	AAY97068	AAB66663	AAY43525	AAB35266	5	AAY92260	4	8	മ	667	AAB35261	693	922	922	922	AAY92256
encoded	4	⊆	racellular	u	ptide seque	conserved	erved	Recombinant interl	trace	Variant Il-1 cytok	\mathbf{r}		Recombinant human	dide	e IL-1ra pol	ra.	IL-1 1	н	enco	_	eukin-IL1	TANGO	æ	۲.		int	ion relate	IL-IL	IL-1 recep	human	c huma	n IL-1	Human IL-1 homolog

ALIGNMENTS

RESULT AAY28408

AAY28408 standard; Protein;

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Human interleukin 1 delta

28-SEP-1999 AAY28408;

(first entry)

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A PRINCE 
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Murine and Human interleukin 1 delta DNA, polypeptides and fragments, useful as molecular weight markers
                                                                                                                                                     WPI; 1999-458310/38.
N-PSDB; AAX89432.
                                                                                                                                                                                                                                                                                                     Sims JE;
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09-JAN-1998;
                                                                                                                                                                                                                                                                                                                                                                                             (IMMV ) IMMUNEX CORP
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98US-0071074
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Best Local
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                                                                                                                                                23-DEC-1998;
22-JAN-1999;
13-APR-1999;
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                         Goddard A,
                                                                                                                                                                                                                                                                             22-DEC-1999;
                                                                                                                                                                                                                                                                                                                                         06-JUL-2000
                                                                                                                                                                                                                                                                                                                                                                                                   WO200039297-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      dermatalogical;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                hIL-1Ra3; human interleukin-1 receptor antagonist-3; IL-11p; osteopathic; interleukin-1-like polypeptide; anti-inflammatory; anti-asthmatic; anti-arthritic; anti-increbial; respiratory; anti-ischemic; vaccine;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human IL-1 receptor antagonist
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          31-OCT-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAY96936
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     weight of a sample protein. The protein and its fragments are useful as controls for peptide fragmentation. This is useful for determining the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The present sequence represents human interleukin 1 delta (:IL-1 delta proteins are useful for the determination of the
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155; Conserv
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99US-0116843.
99US-0129122.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          immunomodulatory; gastrointestinal;
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Pred. No. 1
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RESULT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CC An isolated nucleic acid molecule encoding an interleukin-1-like CC polypeptide (IL-Ilp) that retains one or more activities of the peptide CC from which it is derived, such as the IL-IBR binding activity of a human CC interleukin-1 receptor antagonist-1 (hIL-IRal) polypeptide, is new. The CC nucleic acids may be used in molecular engineering applications, e.g. CC hybridization assays and chromosome and gene mapping studies, for CC recombinantly producing the IL-Ilp polypeptide or for producing gene CC thock out animals to study the role of the protein in metabolism and CC disease processes (conversely, gene therapy protocols may be used to supplement a patients production of the polypeptide or to rectify CC mutations that lead to the production of in active peptides). The CC peptides produced may be used to screen for and produce modulators (e.g. CC intibodies) of IL-Ilp protein expression and activity which may be used CC treat disorders associated with inappropriate IL-Ilp expression and activity, such as inflammatory disorders, asthma, arthritis, costeoarthritis, sepsis, acute lung injury, adult respiratory distress comparate years and continue in the continue production of inflammatory disease.
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                                                                                                                                                                                            Generic; interleukin-1; IL-1; homologue; zilla3; anti-inflammatory; antiagonist; pro-inflammatory; agonist; immunomodulator; antiarthritic; antirheumatic; osteopathic; antipsoriatic; antibacterial; cytostatic; immunosuppressive; antiulcer; antidiabetic; nephrotropic; vasotropic;
                                                                                                               Key
                                                                                                                                                                                                                                                                                                                                                                               AAY92257 standard; Protein; 155
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Nucleic acids encoding interleukin-1-like polypeptides, useful for preventing and treating e.g. inflammation, asthma and psoriasis {\sf prevention}
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2000-452395/39.
N-PSDB; AAA51597.
Peptide
                              Peptide
                                                               Peptide
                                                                                                                                                Homo sapiens
                                                                                                                                                                                                                                                                               Human IL-1 homologue, hzilla3.
                                                                                                                                                                                                                                                                                                                 10-AUG-2000
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                                                                                                Peptide
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14..16
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17..21
                                                                                                               Location/Qualifiers
             'label= beta-strand
                                                                              beta-strand
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Pred. No. 1.2e-86;
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This shows an interleukin-1 (IL-1) homologue, designated zilla: It is believed that zilla3 acts through IL-1 receptors. In ger zilla3 proteins having a Lys residue at position 148 will have anti-inflammatory activity (e.g. AAY92256), whilst those having
                                                                                                                                    Proteins useful for treatment of inflammatory conditions such as rheumatoid arthritis and psoriasis are agonists or antagonists from interleukin-1 homologue
                                                                                                     Disclosure;
                                                                                                                                                                                                                   WPI; 2000-303780/26.
N-PSDB; AAA09193, AA
                                                                                                                                                                                                                                                                                                                 (ZYMO ) ZYMOGENETICS INC
                                                                                                                                                                                                                                                                                                                                                          08-OCT-1998;
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Matches 155
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                                                                                                                                                                                                                                                                                                                                TANGO-93; cytokine; human; secreted protein; II-1 expression; cancer; Interleukin-1 receptor antagonist; II-1ra; inflammation; antiasthmatic; immunosuppressive; antirheumatic; antiarthritic; antipsoriatic; asthma; antiinflammatory; antibacterial; antiulcer; cytostatic; immunomodulator; osteopathic; dermatological; antidiabetic; psoriasis; ulcerative colitis; graft vs-host disease; rheumatoid arthritis; inflammatory bowel disease; septic shock; cachexia; Crohn's disease; chronic myelogenous leukaemia; liver disease; diabetes; osteopathit; Hodgkin's disease; Lyme disease; liver disease; diabetes; osteopathit; Hodgkin's disease; Lyme disease;
                                                                                                                                                                                 06-AUG-1999;
                                                                                                                                                                                                                                                                                                        chromosome 2; systemic
                                                                                                                                                                                                                                                                                                                      autoimmune disease; myasthenia gravis; pharmacogenomic;
                                                                                                                                                    07-AUG-1998;
                                                                                                                                                                                                                17-FEB-2000
                                                                                                                                                                                                                                              WO200008045-A2
                                                                                                                                                                                                                                                                         Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human TANGO-93 protein
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        standard; Protein;
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smatches 0;
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WPI; 2000-205669/18. N-PSDB; AAZ50812.

Isolated nucleic acid sequences encoding TANGO-93 polypeptide useful for treating a variety of cellular processes e.g. asthma, rheumatoid

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18-FEB-2000;
18-FEB-2000;
22-FEB-2000;
01-MAR-2000;
03-MAR-2000;
25-APR-2000;
                                                                                                         01-SEP-1999;
15-SEP-1999;
07-DEC-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      expression. TANGO-93 is useful as a modulating agent for regulating cellular processes like asthma, graft vs-host disease, rheumatoid arthritis, psoriasis, inflammatory bowel disease, septic shock, ulcerative colitis, Crohn's disease, chronic myelogenous leukaemia, cancer, liver disease, Hodgkin's disease, osteoarthritis, Lyme disease, cachexia, and autoimmune diseases e.g. myasthenia gravis, autoimmune diseases e.g. myasthenia gravis, autoimmune diseases are useful in forensic biology, for diagnostic and prognostic assays, prophylactic and therapeutic treatment and pharmacogenomics. The DNA sequences are useful as hybridisation probes and primers, for isolation of TANGO-93 sequence and for the creation of transgenic animals.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The present sequence is the human TANGO-93, a secreted protein that belongs to the cytokine superfamily. It plays a role similar to secreted Interleukin-1 receptor antagonist (ILL-Ira) and its expression is developmentally regulated in the uterus, placent and skeletal muscles. Human TANGO-93 gene is mapped to chromosome 2 within the IL-1 cluster. TANGO-93 modulates immune mediated inflammation and IL-1 gene or protein
                                                                               09-DEC-1999;
11-JAN-2000;
                                                                                                                                                                                       08-MAR-2001
                                                                                                                                                                                                                 WO200116318-A2
                                                                                                                                                                                                                                          Homo sapiens
                                                                                                                                                                                                                                                                    Human; PRO protein; mapping
                                                                                                                                                                                                                                                                                              Human PRO4342.
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 99US-0170262.
2000US-0175481.
2000WO-US04341.
2000WO-US04342.
2000WO-US04414.
2000WO-US05601.
2000US-0187202.
2000US-019397.
                                                                                                                                                             2000WO-US23328
                                                                                                        99WO-US20111.
99WO-US21090.
99US-0169495.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The present sequence is a human PRO polypeptide (secreted and transmembrane). The PRO protein, and PRO agonists, PRO antagonists or anti-PRO antibodies are useful for preparation of a medicament useful in the treatment of a condition which is responsive to the PRO protein, agonists, antagonists or anti-PRO antibodies. The PRO protein may also be employed as molecular weight markers for protein electrophoresis. The PRO coding sequence has applications in molecular biology, including use as hybridisation probes, and in chromosome and gene mapping.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Eaton DL, Fi
Grimaldi CJ,
                                                                                                                                                                                Human; IL-III; interleukin-1 locus; IL-1beta; IL-1receptor; psoriasis; chromosome 2q13; inflammatory disease; heart disease; Graves' disease; rheumatoid arthritis; inflammatory bowel disorder; diabetes; cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Claim 12; Fig 152; 278pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Eighty four nucleic acids encoding molecular biology, including use a chromosome and gene mapping. -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            22-MAY-2000; 2000WO-US14042.
05-JUN-2000; 2000US-0209832.
                                     16-JUL-1999;
                                                                                                                  WO200105974-A2
                                                                                                                                                                   osteoporosis;
                                                                                                                                                                                                                                     Human IL-IL1.
                                                                                                                                                                                                                                                                 08-MAY-2001
                                                                                                                                                                                                                                                                                                                    AAB35260 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence
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                                                             17-JUL-2000; 2000WO-US19508
                                                                                                                                            Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                              LCTVPEADQPVRLTQLPENGGWNAPITDFYFQQCD 155
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DB; AAF92133.
                                                                                                                                                                                                                                                                                                                                                                                lctvpeadqpvrltqlpenggwnapitdfyfqqcd
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J, Gurney AL,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              155 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          . 100.0%; ilarity 100.0%; Conservative
                                                                                                                                                                                                                                                                 (first entry)
                                                                                                                                                                     systemic lupus erythematosus
                                     99US-0144298
                                                                                                                                                                                                                                                                                                                    Protein;
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Watanabe
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Pred. No. 1.2e-86;
0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRO polypeptides, useful in s hybridization probes, and in
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(INTE-) INTERLEUKIN GENETICS INC.

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RESULT
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Best Local Similarity
Matches 155; Conser
          Nucleic acids encoding human and murine interleukin-lLl polypeptides useful for controlling inflammatory processes - \,
                                                                                                                                                                                                                                                                                             Mouse; IL-ILI; interleukin-1 locus; IL-lbeta; IL-lreceptor; psoriasi chromosome 2q13; inflammatory disease; heart disease; Graves' disease rheumatoid arthritis; inflammatory bowel disorder; diabetes; cancer; osteoporosis; systemic lupus erythematosus; human.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The present invention provides the protein and coding sequences of the human and murine interleukin-IL1 (IL-IL1) proteins. The IL-IL1 gene is located between the IL-lbeta and IL-lreceptor genes at human chromosome 2q13. The sequences are useful in the diagnosis, prevention and treatment of heart disease, cancer and inflammatory diseases such as rheumatoid arthritis, systemic lupus erythematosus, inflammatory bowel disorder, diabetes, psoriasis, osteoporosis, lichen sclerosis, ulcerative colitis, severe periodontal disease and pregnancy complications. The present sequence is the human IL-IL1 protein.
                                                                                                                                                                                                                                        WO200105974-A2
                                                                                                                                                                                                                                                                    Unidentified
                                                                                                                                                16-JUL-1999;
                                                                                                                                                                           17-JUL-2000; 2000WO-US19508
                                                                                                                                                                                                                                                                                                                                                                                                        08-MAY-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAB35262 standard;
                                                                                                                   (INTE-) INTERLEUKIN GENETICS
                                                                                                                                                                                                            25-JAN-2001
                                                                                                                                                                                                                                                                                                                                                                       Interleukin-IL1 recombinant protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 11; Fig 3; 150pp; English.
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                                                         2001-091974/10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               VQGGSQCLSCGVGQEPTLTLEPVNIMELYLGAKESKSFTFYRRDMGLTSSFESAAYPGWF 120
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ids encoding human and murine interleukin-1L1 polypeptides
controlling inflammatory processes -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     100.0%; ilarity 100.0%; Conservative 0;
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                                                                                                                                                                                                                                                                                                                                                                                                      (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Protein;
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Pred. No. 1.2e-86;
; Mismatches 0;
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                                                                                                                                                                                                                                                                                                                          IL-lreceptor; psoriasis;
disease; Graves' disease;
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Best Local
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                                                                                                                                                                                                        07-JUL-1999;
13-OCT-1999;
08-DEC-1999;
10-MAR-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The present invention provides the protein and coding sequences of the human and murine interleukin II.1 (II-II) proteins. The II-II.1 gene is located between the II-lbeta and II.1receptor genes at human chromosome 2q13. The sequences are useful in the diagnosis, prevention and treatment of heart disease, cancer and inflammatory diseases such as rheumatoid arthritis, systemic lupus erythematosus, inflammatory bowel disorder, diabetes, psoriasis, osteoporosis, lichen sclerosis, ulcerative colitis, severe periodontal disease and pregnancy complications. The present
                                 Claim 1; Fig 6; 179pp;
                                                        Isolated nucleic acids encoding interleukin-1 (IL-1) antagonist proteins (referred as IL-1Hy1), useful in cancer, e.g. broast adenocarcinoma and brain tumors, inflammatory disease mediated by IL-18
                                                                                                                                              Ford
                                                                                                                      WPI; 2001-071582/08
                                                                                                                                                                                                                                                                     07-JUL-2000; 2000WO-US18710
                                                                                                                                                                                                                                                                                                                       WO200102571-A2
                                                                                                                                                                                                                                                                                                                                                                                            Protein encoded by extended B2HFLS20W cDNA library sequence #2
                                                                                                                                                                                                                                                                                                                                                                                                                                                AAB66664;
                                                                                                                                                                       (HYSE-) HYSEQ
                                                                                                                                                                                                                                                                                                11-JAN-2001
                                                                                                                                                                                                                                                                                                                                                                    interleukin; IL-1 receptor; cancer; inflammation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAB66664 standard;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      sequence is a recombinant IL-IL1 protein
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99US-0457626.
2000US-0523552.
2000US-0576008.
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                                                                                                                                                                       INC.
                                                                                                                                                                                                                                             99US-0348942
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                                 English.
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                                                                    the treatment and an
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The present invention relates to interleukin (IL)-1 antagonist proteins. IL-1Hyl is useful for treating

cancer, receptor

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RESULT
AAY43526
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Best Local
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                                                                                                                                                                 03-APR-1998;
15-MAY-1998;
20-MAY-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              an inflammatory disease mediated by IL-18, inflammation resulting from infection or allergic reactions, and inflammation associated with chronic bronchitis, arthritis, diabetes or
New isolated interleukin-1 receptor binding polypeptides, used
                        N-PSDB;
                                                           Drmanac R,
Leshkowitz
                                                                                                                               31-JUL-1998;
13-JAN-1999;
                                                                                                                                                                                                               05-APR-1999;
                                                                                                                                                                                                                                       14-OCT-1999.
                                                                                                                                                                                                                                                             W09951744-A2
                                                                                                                                                                                                                                                                                                         anti-inflammatory
                                                                                                                                                                                                                                                                                                                   hematopoiesis regulating activity; tissue growth activity; activin activity; inhibin activity; chemotactic activity; chemotactic activity; chemokinetic activity; hemostatic activity; thrombolytic activity;
                                                                                                                                                                                                                                                                                                                                                     graft versus host disease; inflammatory bowel disease; inflammation; pulmonary disease; autoimmune disease; inflammatory disease; antiproliferative; myelogenous leukemia; premature labor; intrauterine infection; nutritional activity;
                                                                                                                                                                                                                                                                                                                                                                                                    Human; interleukin-1 receptor; IL-1; antagonist; sepsis; acute pancreatitis; endotoxic shock; cytokine induced shock; rheumatoid arthritis; chronic inflammatory arthritis; pancreatic cell damage; diabetes mellitus type 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                               A human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      26-JAN-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAY43526 standard; Protein;
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                                                                                            (HYSE-)
                                                                                                                                                       19-JUN-1998;
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                                                        Crkvenjakov
), Kita D,
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                                                                                             INC.
                                                                                                                  98US-0099818.
98US-0127698.
99US-0229591.
99US-0251370.
                                                                                                                                                                 98US-0055010.
98US-0079909.
98US-0082364.
                                                                                                                                                                                                               99WO-US04291
                                                                                                                                                                                                                                                                                                         activity.
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                                                           R, I
                                                         Dickson M,
d J, Pace
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Pred. No. 1.2e-86;
; Mismatches 0;
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                                                         Drmanac S,
, Alfenito
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AAB35263
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    arthritis, chronic inflammatory arthritis, pancreatic cell damage from diabetes mellitus type 1, graft versus host disease, inflammatory bowel disease, inflammatory associated with pulmonary disease, other autolmmune disease, or inflammatory disease, an antiproliferative agent such as for acute or chronic myelogenous leukemia or in the prevention of premature labor secondary to intrauterine infections. They can also exhibit activities such as e.g. nutrificantla activity, cytokine and cell proliferation/differentiation activity, immune stimulating or suppressing activity, hemmatopoiesis regulating activity, tissue growth activity, activity/hibbin activity, chemotactic/chemokinetic activity, activity, activity, activity, activity, chemotactic/chemokinetic activity,
                                                                                                                                                                                                                                                                     Mouse; IL-IL1; interleukin-1 locus; IL-lbeta; IL-lreceptor; psoriasi chromosome 2q13; inflammatory disease; heart disease; Graves' disease rheumatoid arthritis; inflammatory bowel disorder; diabetes; cancer; osteoporosis; systemic lupus erythematosus; human.
               Nicklin M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                hemostatic and thrombolytic activity, receptor/ligand activity, and anti-inflammatory activity. The products can also be used for detection, diagnosis and drug screening.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         receptors (IL-1Rs). The polynucleotides and polypeptides can be use the prevention or treatment of disorders involving sepsis, acute pancreatitis, endotoxic shock, cytokine induced shock, rheumatoid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The present sequence represents a human interleukin-1 (IL-1) receptor antagonist. The polypeptide is capable of binding I
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      e.g. sepsis, shock, arthritis, pancreatitis, graft-versus-host disease, inflammatory disease, autoimmune disease or proliferative disease
                                                 (INTE-) INTERLEUKIN GENETICS INC
                                                                                                                          17-JUL-2000; 2000WO-US19508
                                                                                                                                                                                                  WO200105974-A2
                                                                                                                                                                                                                                      Unidentified.
                                                                                                                                                                                                                                                                                                                                                              Interleukin-IL1 recombinant protein #2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAB35263 standard; Protein; 154
                                                                                     16-JUL-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             121 LCTVPEADQPVRLTQLPENGGWNAPITDFYFQQCD 155
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         61
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ш
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             VQGGSQCLSCGVGQEPTLTLEPVNIMELYLGAKESKSFTFYRRDMGLTSSFESAAYPGWF 120
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         mvlsgalcfrmkdsalkvlylhnnqllagglhagkvikgeeisvvpnrwldaslspvilg
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    155 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    6; 123pp;
                                                                                                                                                                                                                                                                                                                                                                                                   (first entry)
                                                                                     99US-0144298
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         99.6%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 820; DB 20; Pred. No. 2.7e-86;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mismatches
                                                                                                                                                                                                                                                                                                           IL-lreceptor; psoriasis;
lisease; Graves' disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length 155;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               can be used
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                IL-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0,
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Matches 154;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The present invention provides the protein and coding sequences of the human and murine interleukin-ILI (IL-ILI) proteins. The IL-ILI gene is located between the IL-lbeta and IL-lreceptor genes at human chromosome 2q13. The sequences are useful in the diagnosis, prevention and treatment of heart disease, cancer and inflammatory diseases such as rheumatoid arthritis, systemic lupus erythematosus, inflammatory bowel disorder, diabetes, psoriasis, osteoporosis, lichen sclerosis, ulcerative colitis, severe periodontal disease and pregnancy complications. The present sequence is a recombinant IL-ILI protein.
                                                                                                                                                                                                                                                                                            Mouse; IL-ILL; interleukin-1 locus; IL-lbeta; IL-lreceptor; psoriasis; chromosome 2q13; inflammatory disease; heart disease; Graves' disease; rheumatoid arthritis; inflammatory bowel disorder; diabetes; cancer; osteoporosis; systemic lupus erythematosus; human.
                                                                                                      Nicklin M,
                                                                                                                                                                                                                                                                    Unidentified.
            Examples;
                                    Nucleic acids encoding useful for controlling
                                                                                                                                                                                                                                         WO200105974-A2
                                                                                                                                                                                                                                                                                                                                                                                           08-MAY-2001
                                                                                                                                                                                                                                                                                                                                                                                                                    AAB35264
                                                                                                                                                                                                                                                                                                                                                                                                                                              AAB35264 standard; Protein;
                                                                                                                               (INTE-) INTERLEUKIN GENETICS
                                                                                                                                                            16-JUL-1999;
                                                                                                                                                                                   17-JUL-2000; 2000WO-US19508
                                                                                                                                                                                                                25-JAN-2001
                                                                                                                                                                                                                                                                                                                                                               Interleukin-IL1 recombinant protein #3.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Examples;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 121
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         122 CTVPEADQPVRLTQLPENGGWNAPITDFYFQQCD 155
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2 VLSGALCFRMKDSALKVLYLHNNQLLAGGLHAGKVIKGEEISVVPNRWLDASLSPVILGV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2001-091974/10.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        vlsgalcfrmkdsalkvlylhnnqllagglhagkvikgeeisvvpnrwldaslspvilgv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2001-091974/10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      acids encoding human and murine interleukin-1L1 polypeptides
for controlling inflammatory processes -
            Fig
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Fig 6; 150pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 154 AA;
                                                                                                        Barton J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
                                                                                                                                                                                                                                                                                                                                                                                          (first entry)
            <u>ن</u>
                                                                                                                                                            99US-0144298.
         150pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          99.4%; Score 818; DB 22; 100.0%; Pred. No. 4.5e-86;
                                   human and murine interleukin-1L1 polypeptides inflammatory processes -
            English
                                                                                                                                                                                                                                                                                                                                                                                                                                                 157
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0; Mismatches
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Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The present invention provides the protein and coding sequences of the human and murine interleukin-ILI (IL-ILI) proteins. The IL-ILI gene is located between the IL-lbeta and IL-lreceptor genes at human chromosome 2q13. The sequences are useful in the diagnosis, prevention and treatment of heart disease, cancer and inflammatory diseases such as rheumatoid arthritis, systemic lupus erythematosus, inflammatory bowel disorder, diabetes, psoriasis, osteoporosis, lichen sclerosis, ulcerative colitis, severe periodontal disease and pregnancy complications. The present
This polypeptide is variant interleukin-1 (IL-1) homologue, designated zilla3-K148. It is believed that zilla3 acts through IL-1 receptors. In general, zilla3 proteins having a Lys residue at position 148 will have anti-inflammatory activity, whilst those having Asp (see AAY92254).
                                                                                                                                                                                                                                                                                                                                                                                                                                               antagonist; pro-inflammatory; agonist; immunomodulator; antiarthritic;
antirheumatic; osteopathic; antipsoriatic; antibacterial; cytostatic;
immunosuppressive; antiulcer; antidiabetic; nephrotropic; vasotropic;
                                                                             Claim
                                                                                                                        Proteins useful for treatment of inflammatory conditions such as rheumatoid arthritis and psoriasis are agonists or antagonists forms
                                                                                                                                                                                                                  Sheppard PO,
                                                                                                                                                                                                                                                                               08-OCT-1998;
                                                                                                                                                                                                                                                                                                            08-OCT-1999;
                                                                                                                                                                                                                                                                                                                                          13-APR-2000
                                                                                                                                                                                                                                                                                                                                                                         WO200020595-A1
                                                                                                                                                                                                                                                                                                                                                                                                       Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                    vuinerary.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Generic; interleukin-1; IL-1; homologue; zilla3; anti-inflammatory;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human IL-1 homologue, hzilla3-K148
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAY92256;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAY92256 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               sequence is a recombinant IL-IL1 protein.
                                                                                                         new interleukin-1 homologue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           10-AUG-2000 (first entry)
                                                                                                                                                                                                                                               (ZYMO ) ZYMOGENETICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     124
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                64
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      12
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          4 vlsgalcfrmkdsalkvlylhnnqllagglhagkvikgeeisvvpnrwldaslspvilgv 63
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2 VLSGALCFRMKDSALKVLYLHNNQLLAGGLHAGKVIKGEEISVVPNRWLDASLSPVILGV
                                                                                                                                                                                   2000-303780/26
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               qggsqclscgvgqeptltlepvnimelylgakesksftfyrrdmgltssfesaaypgwfl
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       154;
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                                                                                                                                                                      AAA09195
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                                                                           56-57; 64pp;
                                                                                                                                                                                                                  West RR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AA;
                                                                                                                                                                                                                                                                               98US-0169745
                                                                                                                                                                                                                                                                                                            99WO-US23533
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      99.4%;
                                                                                                                                                                                                                  Clegg
                                                                           English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          155
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 818; DB 22;
Pred. No. 4.6e-86;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                123
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RESULT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Zilla3 is used to modulate an immune response in an animal (claimed). Antagonists zilla3 forms may be used to treat or prevent chronic inflammatory diseases such as rheumatorid arthritis, osteoarthritis and Lyme arthritis, psoriasis, to reduce tissue damage after ischemia, to treat septic shock, graft-versus-host disease and leukemia. The antagonists may also alleviate inflammatory bowel disease including Crohn's disease and ulcerative colitis, insulin-dependent diabetes mellitus, acute pancreatitis, glomerulonephritis and cerebral ischemia. Agonist forms of zilla3 may promote wound healing by IL-1 effects on growth factor secretion and cell proliferation. They may also treat infertions especially restriction.
Proteins useful for treatment of inflammatory conditions such as rheumatoid arthritis and psoriasis are agonists or antagonists for
                                                                        Sheppard PO,
                                                                                                                                                            08-OCT-1999;
                                                                                                                                                                                                                                                                Misc-difference 99
                                                                                                                                                                                                                                                                                                                                      Ното
                                                                                                                                                                                                                                                                                                                                                                             antagonist; pro'inflammatory; agonist; immunomodulator; antiarthritic;
antirheumatic; osteopathic; antipsoriatic; antibacterial; cytostatic;
immunosuppressive; antiulcer; antidiabetic; nephrotropic; vasotropic;
                                                                                                                                                                                                                                                                                                                                                                                                                         Generic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                     Hhuman IL-1 homologue, hzilla3-D148
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAY92254 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence
                                                                                                   (ZYMO ) ZYMOGENETICS INC
                                                                                                                                                                                         13-APR-2000
                                                                                                                                                                                                                    WO200020595-A1
                                                                                                                                                                                                                                                                                           Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  10-AUG-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         infections, especially gastrointestinal infections.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 121
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  VQGGSQCLSCGVGQEPTLTLEPVNIMELYLGAKESKSFTFYRRDMGLTSSFESAAYPGWF 120
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          LCTVPEADQPVRLTQLPENGGWNAPITDFYFQQCD 155
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   vqggsqclscgvgqeptltlepvnimelylgakesksftfyrrdmgltssfesaaypgwf 120
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       is used
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                         interleukin-1; IL-1; homologue; zilla3; anti-inflammatory;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 155 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (first entry)
                                                                      West RR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       position will have pro-inflammatory action. to modulate an immune response in an animal
                                                                                                                                98US-0169745
                                                                                                                                                            99WO-US23533
                                                                                                                                                                                                                                                                                           Location/Qualifiers
                                                                                                                                                                                                                                                 /label= Ala, Ile, Thr
                                                                                                                                                                                                                                                                           /label= Glu, Lys
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    99.18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 816; DB 21;
Pred. No. 7.6e-86;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length 155;
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   forms
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RESULT 14
AAY92253
ID AAY922
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QΥ
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            chronic inflammatory diseases such as rheumatoid arthritis, costeoarthritis and Lyme arthritis, psoriasis, to reduce tissue damage after ischemia, to treat septic shock, graft-versus-host disease and leukemia. The antagonists may also alleviate inflammatory bowel disease including Crohn's disease and ulcerative collitis, insulin-dependent diabetes mellitus, acute pancreatitis, glomerulonephritis and cerebral ischemia. Agonist forms of zilla3 may promote wound healing by IL-1 effects on growth factor secretion and cell proliferation. They may also treat infections, especially gastrointestinal infections. Note: This sequence is not given in the specification, it is created from SEQ. ID. 7, which is given on pages 55-56.
                                                                                                                                                                                                                                                                                          Generic; interleukin-1; IL-1; homologue; zilla3; anti-inflammatory; antagonist; pro-inflammatory; agonist; immunomodulator; antiarthritic; antirheumatic; osteopathic; antipsoriatic; antibacterial; cytostatic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             This is a generic interleukin-1 (II-1) homologue, designated zilla3. It is believed that zilla3 acts through II-1 receptors. In general, zilla3 proteins having a Lys residue at position 148 (see AAY92255 and AAY92256) will have anti-inflammatory activity, whilst those having Asp (i.e. this sequence) or Glu at this position will have pro-inflammatory action. Zilla3 is used to modulate an immune response in an animal (claimed). Antagonists zilla3 forms may be used to treat or prevent
                                                 WO200020595-A1
                                                                                                      Misc-difference
                                                                                                                                     Misc-difference
                                                                                                                                                                      Misc-difference
                                                                                                                                                                                                                                                                                                                                                                     Generic human
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAY92253 standard; protein; 155 AA
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                                                                                                                                                                                                                            Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             new interleukin-1 homologue
                                                                                                                                                                                                                                                                             mmunosuppressive; antiulcer; antidiabetic; nephrotropic; vasotropic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  121
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  LCTVPEADQPVRLTQLPENGGWNAPITDFYFQQCD 155
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 lctvpeadqpvrltqlpenggwnapitdfyfqqcd
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    VQGGSQCLSCGVGQEPTLTLEPVNIMELYLGAKESKSFTFYRRDMGLTSSFESAAYPGWF 120
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      mvlsgalcfrmkdsalkvlylhnngllagglhagkvikgexisvvpnrwldaslspvilg 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                      (first entry)
                                                                                                                                                                                                                                                                                                                                                                   IL-1 homologue, hzilla3-X148
                                                                                                   /label≔ Ala,
148
                                                                                                                                                                                         Location/Qualifiers
                                                                                  /label=
                                                                                                                                                      /label=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           98.7%;
98.7%;
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                                                                                Glu,
                                                                                                                                                      Glu,
                                                                                  Lys, Asp
                                                                                                                     Ile,
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Pred. No. 2.2e-85;
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Best Local :
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 Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         zilla3. It is believed that zilla3 acts through IL-1 receptors. In general, zilla3 proteins having a Lys residue at position 148 (see AAY92255) will have anti-inflammatory activity, whilst those having Asp (see AAY92254) or Glu at this position will have pro-inflammatory action. Zilla3 is used to modulate an immune response in an animal (claimed). Antagonists zilla3 forms may be used to treat or prevent chronic inflammatory diseases such as rheumatoid arthritis, osteoarthritis and Lyme arthritis, psoriasis, to reduce tissue damage after ischamia, to treat septic shock, graft-versus-host disease and leukemia. The antagonists may also alleviate inflammatory bowel disease including Crohn's disease and ulcerative collits, insulin-dependent disease and alleviate inflammatory bowel disease.
                                                                                                 Generic; interleukin-1; IL-1; homologue; zilla3; anti-inflammatory; antagonist; pro-inflammatory; agonist; immunomodulator; antiarthritic; antiheumatic; osteopathic; antipsoriatic; antibacterial; cytostatic; immunosuppressive; antiulcer; antidiabetic; nephrotropic; vasotropic;
                                                                                                                                                                                           Generic human IL-1 homologue,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          diabetes mellitus, acute pancreatitis, glomerulonephritis and cerebral ischemia. Agonist forms of zilla3 may promote wound healing by IL-1 effects on growth factor secretion and cell proliferation. They may a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Proteins useful for treatment of inflammatory conditions such as rheumatoid arthritis and psoriasis are agonists or antagonists forms of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 1; Page 55-56; 64pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sheppard PO,
                                                   Homo sapiens
                                                                                      vulnerary.
                                                                                                                                                                                                                               10-AUG-2000
                                                                                                                                                                                                                                                                                                  AAY92255 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               treat infections, especially gastrointestinal infections.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  new interleukin-1 homologue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (ZYMO ) ZYMOGENETICS INC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                              VQGGSQCLSCGVGQEPTLTLEPVNIMELYLGAKESKSFTFYRRDMGLTSSFESAAYPGWF 120
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MVLSGALCFRMKDSALKVLYLHNNQLLAGGLHAGKVIKGEEISVVPNRWLDASLSPVILG
                                                                                                                                                                                                                                                                                                                                                                                        lctvpeadqpvrltqlpenggwnapitxfyfqqcd 155
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              \verb|mvlsgalcfrm| kdsalkvlylhnnqllaggl| hagkvlkgexisvvpnrwldaslspvilg|
                                                                                                                                                                                                                                                                                                                                                                                                                                                            vqggsqclscgvgqeptltlepvnimelylgakesksfxfyrrdmgltssfesaaypgwf 120
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            polypeptide is a generic interleukin-1 (IL-1) homologue, designated
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              155
                                                                                                                                                                                                                            (first entry)
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Location/Qualifiers
                                                                                                                                                                                                                                                                                                  protein; 155
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98.1%;
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                                                                                                                                                                                             hzilla3-K148
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 155;
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Search completed: Job time: 27 sec

June

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2001, 14:05:25

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CC This is a generic interleukin-1 (IL-1) homologue, designated zilla3. CC It is believed that zilla3 acts through IL-1 receptors. In general, CC zilla3 proteins having a Lys residue at position 188 (this sequence CC and AAY92256) will have anti-inflammatory activity, whilst those having Asp (AAY92254) or Glu at this position will have pro-inflammatory CC action. Zilla3 is used to modulate an immune response in an animal CC (claimed). Antagonists zilla3 forms may be used to treat or prevent CC chronic inflammatory diseases such as rheumatoid arthritis, CC osteoarthritis and Lyme arthritis, psoriasis, to reduce tissue damage CC after ischemia, to treat septic shock, graft-versus-host disease and CC leukemia. The antagonists may also alleviate inflammatory bowel disease including Crohn's disease and ulcerative colitis, insulin-dependent CC diabetes mellitus, acute pancreatitis, glomerulonephritis and cerebral ischemia. Agonist forms of zilla3 may promote wound healing by IL-1 CC effects on growth factor secretion and cell proliferation. They may CC also treat infections, especially gastrointestinal infections.

CC Note: This sequence is not given in the specification, it is created. CC from SEQ. ID. 7, which is given on pages 55-56.
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                                                                                                                                                                                                                                                                                  Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 3; Page -; 64pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sheppard PO,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              08-OCT-1999;
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   121
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                                                                                                                                                                                                                Local Similarity
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                                                                                                                                             1 MVLSGALCFRMKDSALKVLYLHNNQLLAGGLHAGKVIKGEEISVVPNRWLDASLSPVILG
mvlsgalcfrmkdsalkvlylhnnqllagglhagkvikgexisvvpnrwldaslspvilg
                                LCTVPEADQPVRLTQLPENGGWNAPITDFYFQQCD 155
                                                                                                                                                                                                                                                                                  155 AA;
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99
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                                                                                                                                                                                                              97.8%;
98.1%;
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                                                                                                                                                                                                              Score 805; DB 21;
Pred. No. 1.4e-84;
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Result
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Maximum
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Perfect score:
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length: 2000000000
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1: pir1:*
2: pir2:*
3: pir3:*
4: pir4:*
219241 segs, 76174552 residues
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Gapop 10.0 ,
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823
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MVLSGALCFRMKDSALKVLY.....LPENGGWNAPITDFYFQQCD
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pir2:*
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interleukin-1 beta
interleukin-1 chic
                                                                      methionyl-tRNA syn
methionyl-tRNA syn
probable SNF2-type
hypothetical prote
                                           hypothetical prote phosphotransferase
                                                                                                                                                                            ATP sulfurylase -
polyprotein - marm
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H+-transporting AT
N-methylhydantoina
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45	44	43	42	41	40	39	38	37	36	35	34	33	32	31	30
69	69.5	70	70	70	70.5	70.5	71	71	71	71.5	71.5	71.5	71.5	71.5	. 72
8.4	8.4	8.5	8.5	8.5	8.6	8.6	8.6	8.6	8.6	8.7	8.7	8.7	8.7	8.7	8.7
152	801	323	305	269	1161	778	611	535	475	1110	1036	700	578	551	423
2	N	N	N	N	N	N	N	2	N	N	N	N	Ν	N	N
B43863	в86673	A82433	E83091	C83516	T18642	D72421	A48582	T06285	G84697	T33877	B83466	T34321	T45107	H72117	B86214
hypothetical prote	penicillin-binding	fructokinase VCA06		hypothetical prote	hypothetical prote	xylosidase - Therm	vacuolar ATPase A			hypothetical prote	probable RND efflu	hypothetical prote	H+-transporting AT	methioninetRNA l	hypothetical prote

ALIGNMENTS

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interleukin-1 receptor antagonist secreted form precursor - human (:Species: Homo sapiens (man) (:Date: 07-Jun-1990 #sequence_revision 07-Jun-1990 #text_change 26-May-2000 (:Accession: A40956; I37894; A30368; S08160; S08159; A37822 R;Eisenberg, S.P.; Brewer, M.T.; Verderber, E.; Heimdal, P.; Brandhuber, B.J.; The Proc. Natl. Acad. Sci. U.S.A. 88, 5232-5236, 1991 A;Title: Interleukin 1 receptor antagonist is a member of the interleukin 1 gene: A. Reference number: A40956; MUID:91271363 A;Accession: A40956 MUID:91271363 A;Accession: A40956 Accession: A40956 MUID:91271363 A;Accession: A40956 MUID:91271363 A;Cross-references: GB:M63099; NID:9186385; PIDN:AAB41943.1; PID:9186386 A;Cross-references: GB:M63099; NID:9186385; PIDN:AAB41943.1; PID:9186385; PIDN:PID:9186385; PIDN:PID:9186385; PIDN:PID:918
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A;Title: ILIHY1: A novel interleukin-1 receptor antagonist gene.
A;Reference number: JC7104; MUID:99443727
A;Accession: JC7104
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c;SpecLes: Homo sapiens (man)
c;Date: 03-Dec.1999 #sequence_revision 03-Dec-1999 #text_change 21-Jul-2000
C;Accession: JC7104
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A30368
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A;Map position: 2ql4
C;Keywords: macrophage
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A; Residues: 1-155 < MUL>
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        MVLSGALCFRMKDSALKVLYLHNNQLLAGGLHAGKVIKGEEISVVPNRWLDASLSPVILG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              155;
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Pred. No. 2.3e-77;
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                                                                                                                                                                                                                                                                                                                                                       Brandhuber, B.J.; Thomps
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C;Keywords: alternative splicing; cytokine receptor; extracellular protei
F;1-25/Domain: signal sequence #status predicted <SIG>
F;26-177/Product: interleukin-1 receptor antagonist #status experimental
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              A;Molecule type: protein
A;Residues: 26-52;70-77:122-127;170-175 <BIE>
A;Experimental source: culture medium, PMA-stimulated THP-1
C;Comment: For an alternative splice form, see PIR:A39386
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      A; Molecule type: protein
A; Residues: 26-75; 97-108;110-116;120-131;163-176 <HAN>
R; Bienkowski, M.J; Eessalu, T.E.; Berger, A.E.; Trueso
J. Biol. Chem. 265, 14505-14511, 1990
A; Title: Purification and characterization of interleul
A; Reference number: A37822; MUID:90354444
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            A;Cross-references: GB:X52015; NID:g32576; R;Hannum, C.H.; Wilcox, C.J.; Arend, W.P.; Nature 343, 336-340, 1990
A;Title: Interleukin-1 receptor antagonist A;Reference number: S08159; MUID:90136920
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   A;Cross-references: EMBL:X64532; NID:g33798; PIDN:CAA45832.1; R;Carter, D.B.; Deibel Jr., M.R.; Dunn, C.J.; Tomich, C.S.C.; J.G.; Sieu, L.C.; Hardee, M.M.; Zurcher-Neely, H.A.; Reardon, Nature 344, 633-638, 1990
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A; Map position: 2914.2-2914.2
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A; Residues: 1-177 <LEN>
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A; Accession: I37894
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      A; Molecule type: mRNA
A; Residues: 1-177 <EI2>
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RESULT
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Best Local 9
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                                                                                                                                                                                                                                                                                                                                                                    Matches
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                                                                                                                     126 EADQPVRLTQLPENGGWNAPITDFYFQQ 153
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                                                                                                                                                                     CLSCVKSGDETRLQLEAVNITDLSENRKQDKRFAFIRSDSGPTTSFESAACPGWFLCTAM 150
                                                                                                                                                                                                                CLSC-GVGQEPTLTLEPVNIMELYLGAKESKSFTFYRRDMGLTSSFESAAYPGWFLCTVP 125
                                                                                                                                                                                                                                                                      FRIWDVNQKTFYLRNNQLVAGYLQGPNVNLEEKIDVVP-----
                                                                                                                                                                                                                                                                                                         FRMKDSALKVLYLHNNQLLAGGLHAGKVIKGEEISVVPNRWLDASLSP--VILGVQGGSQ
                                                                       EADQPVSLTNMPDEG - - - VMVTKFYFQE
                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             A30368;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          cloning, expression and biological characterization 0368; \mathtt{MUID}: 90220867
                                                                                                                                                                                                                                                                                                                                                                                      38.8%;
                                                                                                                                                                                                                                                                                                                                                                    13;
                                                                                                                                                                                                                                                                                                                                                               Score 319.5;
Pred. No. 1.7e
13; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PIDN:CAA36262.1; PID:g32577 Joslin, F.G.; Dripps, D.J.;
                                                                                                                                                                                                                                                                                                                                                                                   1.7e-25;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Truesdell,
                                                                                                                                                                                                                                                                                                                                                                    49;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           extracellular protein; glycoprotein
                                                                                                                                                                                                                                                                   -IEPHALFLGIHGGKM 90
                                                                                                                                                                                                                                                                                                                                                                    Indels
                                                                                                                                                                                                                                                                                                                                                                                                                  Length 177;
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the mature protein, were con
; Brewer, M.T.; Hannum, C.H.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  receptor level
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PID:g33799
Laborde, A.L.; Sl
I.M.; Heinrikson,
                                                                                                                                                                                                                                                                                                                                                                 13;
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son, R.L.;
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A:Map position: 2q14.2-2q14.2 (Superfamily: interleukin-1 (C; Superfamily: interleukin-1 (C; Keywords: alternative splicing; cytokine receptor F;1-180/Product: interleukin-1 receptor antagonist, long F;1-3,25-180/Product: interleukin-1 receptor antagonist,
                                                                                                                                                                                                                                                                                                                                                                 A; Molecule type: mRNA
A; Residues: 1-3,25-180 < HAS>
A; Cross-references: GB: M55646; NID: 9186291;
C; Comment: For an alternative splice form, s
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                R:Haskill, S.; Martin, G.; Van Le, L.; Morris, J.; Peace, A.; Bigler, C.F.; Jaffe, Proc. Natl. Acad. Sci. U.S.A. 88, 3681-3685, 1991
A;Title: cDNA cloning of an intracellular form of the human interleukin 1 receptor A;Reference number: A39386; MUID:91219436
A;Accession: A39386
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    N;Contains: interleukin-1 receptor antagonist, short intracellular splice fi C;Species: Homo sapiens (man) C;Date: 28-Feb-1992 #sequence_revision 11-Apr-1997 #text_change 26-May-2000
                                                                                                                                                                                                                                                                                                               A; Gene: GDB: IL1RN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             A;Cross-references: EMBL:X84348; NID:g1008970; PIDN:CAA59087.1; PID:g1008971 R;Haskill, S.; Martin, G.; Van Le, L.; Morris, J.; Peace, A.; Bigler, C.F.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     A; Molecule type: mRNA
A; Residues: 1-180 < RES>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        C;Accession: I37893; A39386 R;Muzio, M.; Polentarutti,
                                                                                                                                                                                                                                                                                    A; Cross-references: GDB:125897;
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                          Query Match
Best Local 9
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                                Similarity
     Conservative
                             38.8%;
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  13;
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  Score 319.5; DB 2
Pred. No. 1.7e-25;
3; Mismatches 49
                                                                                                                                                                                                                                                                                                                                                                       ; PIDN:AAA59138.1; see PIR:A30368
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        49;
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short intracellular splice f
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C;Date: 09-Sep-1994 #sequence_revision 09-Sep-1994 #text_change 16-Jul-1999 C;Accession: A44610; B40956; A49031; I56106; I52970 R;Matsushime, H.; Roussel, M.F.; Matsushima, K.; Hishinuma, A.; Sherr, C.J.
                     A;Cross-references: GB:M64404; NID:g198296; PIDN:AAA39277.1; R;Eisenberg, S.P.; Brewer, M.T.; Verderber, E.; Heimdal, P.; Proc. Natl. Acad. Sci. U.S.A. 88, 5232-5236, 1991
A;Title: Interleukin I receptor antagonist is a member of the A;Reference number: A40956; MUID:91271363
                                                                                                                                                      A; Molecule type: mRNA
A; Residues: 1-178 <MAT>
                                                                                                                                                                                                                            A; TILLE: Cloning and expression of murine interleukin-1 A; Reference number: A44610; MUID:91316273
                                                                                                                                                                                                                                                             Blood 78, 616-623, 1991 A; Title: Cloning and ex
                                                                                                                                                                                                                                                                                                                                                                                             C; Species: Mus musculus (house mouse)
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                                                                                                    Brandhuber,
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                                                   interleukin 1 gene fami
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R; Zahedi, K.; Seldin, M.F.; Rits, M.; Ezekowitz, R.B.; Whitehead, A.S.
J. Immunol. 146, 4228-4233, 1991
A; Title: Mouse IL-1 receptor antagonist protein: Molecular characterization, A; Reference number: 156106; MUID:91250712
A; Accession: 156106
                                                                                                                                  R;Eisenberg, S.P.; Brewer, M.T.; Verderber, E.; Heimdal, P.; Brandhuber, B.J.; Proc. Natl. Acad. Sci. U.S.A. 88, 5232-5236, 1991
A;Title: Interleukin 1 receptor antagonist is a member of the interleukin 1 gen A;Reference number: A40956; MUID:91271363
A;Accession: C40956
                                                                                                                                                                                                                                               C;Species: Rattus norvegicus (Norway ra
C;Date: 20-Mar-1992 #sequence_revision
C;Accession: C40956
                                                                                                                                                                                                                                                                                                                                                   RESULT
C40956
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C;Keywords: cytokine receptor
F;1-26/Domain: signal sequence #status predicted <SIG>
F;27-178/Product: interleukin-1 receptor antagonist #status predicted <MA2>
                     A;Cross-references: GB:M63101;
C;Superfamily: interleukin-1
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A;Accession: I52970
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Cytokine 6, 1-9, 1994
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  C; Keywords:
                                                                   A; Molecule type: DNA
A; Residues: 1-178 <EIS>
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A; Residues: 1-178 <RE2>
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                                                                                                                 A; Status: preliminary
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DQPVRLTQLPENGGWNAP -- ITDFYFQQ 153
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SCG-VGQEPTLTLEPVNIMELYLGAKESKSFTFYRRDMGLTSSFESAAYPGWFLCTVPEA 127
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FRIWDTNQKTFYLRNNQLIAGYLQGPNIKLEEKIDMVP----IDLHSVFLGIHGGKLCL
cytokine receptor
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                                              NID:g204928;
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Pred. No. 2.3e-24;
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                                              PIDN: AAA41434.1;
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                                           PID: g204929
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interleukin-1 receptor antagonist secreted form precursor - rabbit C;Species: Oryctolagus cuniculus (domestic rabbit) C;Date: 06-Oct-1994 #sequence_revision 18-Nov-1994 #text_change 16-Jul-1999 C;Accession: A54377; I46729 R;Cominelli, F; Bortolami, M; Pizarro, T.T.; Monsacchi, L.; Ferretti, M.; P.; Bolol. Chem. 269, 6962-6971, 1994 A;Title: Rabbit interleukin-1 receptor antagonist. Cloning, expression, func A;Reference number: A54377; MOID:94165101 A;Accession: A54377 A;Molecule type: mRNA A;Residues: 1-177 <COM>
interleukin-1 beta precursor - sheep
W;Alternate names: hematopoietin-1; IL-1 beta
C;Species: Ovis orientalis aries, Ovis ammon aries (domestic sheep)
C;Date: 08-Jun-1994 #sequence_revision 22-Nov-1996 #text_change 15-Oct-1999
C;Accession: S23010; S43047; S13092; B61246
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                C;Keywords: cytokine receptor; extracellular protein; F;1-25/Domain: signal sequence #status predicted <SIG>F;109/Binding site: carbohydrate (Asn) (covalent) #status
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           A;Molecule type: mrNA
A;Residues: 1-177 <GGT>
A;Residues: 1-177 <GGT>
A;Cross-references: GB21832; NID:g425787; PIDN:BAA04860.1; PID:g452205
C;Superfamily: interleukin-1
C;Superfamily: interleukin-1
C;Superfamily: fine recentor: extracellular protein; glycoprotein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             RESULT
A54377
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        A; Title: Interleukin-1 recetor antagonist in A; Reference number: I46729; MUID: 93052512
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              A; Note: sequence extracted from NCBI backbook; Goto, F.; Goto, K.; Miyata, T.; Ohkawara, Immunology 77, 235-244, 1992
                                                                                                                    $23010
                                                                                                                                       RESULT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           A;Accession: I46729
A;Status: translated from GB/EMBL/DDBJ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             A; Experimental source: colon tissue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                A;Cross-references: GB:S68977; NID:g545740;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local S
Matches 67
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local :
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                                                                                                                                                                                                                                                                                                     CLSCVKSGDKMKLHLEAVNITDLGKNKEQDKRFTFIRSNSGPTTTFESASCPGWFLCTAL
                                                                                                                                                                                                                                                                                                                                                                                                                                     FRMKDSALKVLYLHNNQLLAGGLHAGKVIKGEEISVVPNRWLDASLSP--VILGVQGGSQ 66
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DHPVSLTNTPKE---PCTVTKFYFQE 176
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DQPVRLTQLPENGGWNAPITDFYFQQ 153
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SCVKSGDDTKLQLEEVNITDLNKNKEEDKRFTFIRSETGPTTSFESLACPGWFLCTTLEA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FRMKDSALKVLYLHNNQLLAGGLHAGKVIKGEEISVVPNRWLDASLSPVILGVQGGSQCL 68
                                                                                                                                                                                                          EADQPVSLTNTPDD---SIVVTKFYFQE
                                                                                                                                                                                                                                                     EADQPVRLTQLPENGGWNAPITDFYFQQ 153
                                                                                                                                                                                                                                                                                                                                               CLSC-GVGQEPTLTLEPVNIMELYLGAKESKSFTFYRRDMGLTSSFESAAYPGWFLCTVP 125
                                                                                                                                                                                                                                                                                                                                                                                          FRIWDVNQKTFYLRNNQLVAGYLQGPNAKLEERIDVVP-----LEPQLLFLGIQRGKL
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FRIWDTNQKTFYLRNNQLIAGYLQGPNTKLEEKIDMVP----IDFRNVFLGIHGGKLCL
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45.98;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     17;
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Pred. No. 4.1
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Mismatches
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interleukin-1 beta precursor - bovine
N;Alternate names: hematopoietin-1; IL-1 beta
C;Species: Bos primigenius taurus (cattle)
C;Date: 31-Mar-1989 #sequence_revision 31-Mar-1989 #text_change 22-Jun-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          A; Molecule type: mRNA, A; Molecule type: mRNA, A; Residues: 1-144, 'I.', 146-266 <AND>
A; Residues: 1-144, 'I.', 146-266 <AND>
C; Comment: This protein lacks a conventional signal sequence for protein export. Cleavag ved form of interleukin-lbeta, unlike interleukin 1-alpha, is inactive.

ved form of interleukin-lbeta, unlike interleukin 1-alpha, is inactive.
                                              C;Accession: JL0010; S01380
R;Maliszewski, C.R.; Baker, P.E.; Schoenborn, M.A.;
MO1. Immunol. 25, 429-437, 1988
A;Title: Cloning, sequence and expression of bovine
A;Reference number: A94695; MUID:88318652
A;Accession: JL0010
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A; Residues: 1-13, 'C', 15-54, 'K', 56-63, 'A', 65-144, 'L', 146-266 <
A; Residues: 1-13, 'C', 15-54, 'K', 56-63, 'A', 65-144, 'L', 146-266 <
A; R: Gross-references: EMBL: X54796; NID: g1273; PIDN: CAA38566.1;
R: Fiskerstrand, C.; Sargan, D.
Nucleic Acids Res. 18, 7165, 1990
A; Title: Nucleotide sequence of ovine interleukin-1 beta.
A; Reference number: S13092; MUID: 91088326
A; Accession: S13092
                       A; Molecule type: mRNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 C;Keywords: cytokine; immunoregulation; inflammation; lymphokine; macrophage; mitogen F;114-266/Product: interleukin-1 beta #status predicted <MAT>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    C; Genetics:
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A;Accession: B61246
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A;Note: the authors translated the codon AGT for residue 62 R;Andrews, A.E.; Barcham, G.J.; Brandon, M.R.; Nash, A.D. Immunology 74, 453-460, 1991
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    A; Molecule type: mRNA
A; Residues: 1-13,'C',15-54,'K',56-61,'S',63,'A',65-144,'L',146-266
A; Cross-references: EMBL: X54796
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A; Note: the sequence from Fig. 1
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A; Residues: 1-266 <SEO>
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A; Accession: S23010
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A;Title: Nucleotide sequence of ovine macrophage interleukin-1 beta
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DNA Seg. 1, 423-426, 1991
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         C; Superfamily: interleukin-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         A;Gene: IL-1-beta
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                                        249
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Match 17.2%;
Local Similarity 31.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TPTLQLEEVD-PKVYPKRNMEKREVEYKTEIKNTVEFESVLYPNWYISTSQIEEKPVFLG
                                                                                                                                                                                                                                                                                                                                                                                                                                                        RF--RGGQD--ITDF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          QLPENGGWNAPITDF 149
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   EPTLTLEPVNIMELYLGAKESKSFTFYRRDMGLTSSFESAAYPGWFLCTVPEADQPVRLT 134
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              LKALHLPSQEMSREVVFCMSFVQGEERD------NKIPVALGIRDKNLYLSCVKKGD 189
1 - 266
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
  <MAL>
                                                                                                                                                                                                                                                                                                                                                                                                                                                     259
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 141.5; DB 1
Pred. No. 5.7e-07;
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                                                                                                         interleukin 1-alpha and interleukin
                                                                                                                                                              Davis, B.S.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length 266;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          15;
                                                                                                                                                              Cosman, D.;
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interleukin-1 beta precursor - pig
C:Species: Sus scrofa domestica (domestic pig)
C:Date: 20-May-1994 #sequence_revision 01-Dec-1995 #text_change 16-Jul-1999
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A; Residues: 1-251, 'A', 253-266 <LEO>
A; Residues: 1-251, 'A', 253-266 <LEO>
A; Cross references: EMBL: X12498; NID: 9448; PIDN: CAA31018.1; PID: 9449
C; Comment: This protein is a cytokine that mediates a variety of immunoregulatory C; Comment: This protein lacks a conventional signal sequence for protein export.
C; Comment: Interleukin-1beta, unlike interleukin 1-alpha, is inactive.
C; Comment: Interleukin-1beta precursor is less heavily myristoylated than interleukin-1c; Superfamily: interleukin-1
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A; Title: Gene sequence, cDNA construction, A; Reference number: S38373; MUID:94039070
A; Accession: S38373
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                                                                                                                                                                                                                                                                                                                                                                                                                                                               C; Genetics:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         A; Molecule type: DNA
A; Residues: 1-267 < VAN>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   A; Cross-references: EMBL: X74568; NID: g407899;
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Best Local Similarity
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Best Local Similarity
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235 YISTSQAEQKPVFL---GNSKGRQDITDF
                                          120 FLCTVPEADQPVRLTQLPENGGWNAPITDF 149
                                                                                                                                                                                         132 LVLAG-----PHELKALHLLKGDLKREVVFCMSFVQGDD-----SDDKIPVTLG 175
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                                                                                                                      75 EPTLTLEPVNIMELYLGAKESKSFTFYRRDMGLTSSFESAAYPGWFLCTVPEADQPVRLT 134
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                                                                                          IKGKNLYLSCVMKDDTPTLQLEDVD-PKSYPKRDMEKRFVFYKTEIKNRVEFESALYPNW
                                                                                                                                                                                                                                       MVLSGALCFRMKDSALKVLYLHNNQLLAGGLHAGKVIKGEEISVVPNRWLDASLSPVILG 60
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                                                                                                                                                                                                                                                                                      15.8%; Score 130; DB 2; ilarity 29.3%; Pred. No. 8.7e-06; Conservative 21; Mismatches 63;
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                                                                                                                                                                                                                                                                                                                                 Length 267
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R; Huether, M.J.; Lin, G.; Smitl
Gene 129, 285-289, 1993
A; Title: Cloning, sequencing ar
A; Reference number: JN0724; MU
A; Accession: JN0724
A; Molecule type: mRNA
A; Molecule type: mRNA
A; Residues: 1-267 < HUE>
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A;Experimental source: alveolar macrophage
C;Comment: This protein is a pleiotropic cytokine that mediates a variety of processes in C;Comment: This protein lacks a conventional signal sequence for protein export. Cleavaged form of interleukin-lacks a conventional signal sequence for protein export. Cleavaged form of interleukin-lacks a conventional signal sequence for protein export. Cleavaged form of interleukin-lacks a conventional signal sequence for protein export. Cleavaged form of interleukin-lacks a conventional signal sequence for protein export. Cleavaged form of interleukin-lacks a conventional signal sequence for protein export. Cleavaged form of interleukin-lacks a conventional signal sequence for protein export. Cleavaged form of interleukin-lacks a conventional signal sequence for protein export. Cleavaged form of interleukin-lacks a conventional signal sequence for protein export. Cleavaged form of interleukin-lacks a conventional signal sequence for protein export. Cleavaged form of interleukin-lacks a conventional signal sequence for protein export. Cleavaged form of interleukin-lacks a conventional signal sequence for protein export. Cleavaged form of interleukin-lacks a conventional signal sequence for protein export.
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Gene 177, 11-16, 1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              C;Species: Equus caballus (domestic horse)
C;Date: 28-Oct-1997 #sequence_revision 28-Oct-1997 #text_change 20-Jun-2000
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C:Date: 14-Jul-1994 #sequence_revision 22-Nov-1996 #text_change 22-Jun-1999
C:Accession: JN0724
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A; Residues: 1-214 <KAT>
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                                                                                                                                                                                                                                                                    Matches
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Best Local Similarity
      158
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                                                                                                                                                                   IKGEEISVVPNRWLDASLSPVILGVQGGSQCLSCGVGQ-EPTLTLEPVNIMELYLGAKES 95
KRFVFNKMEIKGNVEFESAMYPNWYISTSQAEKSPVFLGN--TRGG--RDITDFIME 210
                                                           KSFTFYRRDMGLTSSFESAAYPGWFLCTVPEADQPVRLTQLPENGGWNAPITDFYFQ 152
                                                                                                                                   VQGEE---
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             YISTSQAEQKPVFL----GNSKGRQDITDF 260
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FLCTVPEADQPVRLTQLPENGGWNAPITDF 149
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              IKGKNLYLSCVMKDNTPTLQLEDID-PKRYPKRDMEKRFVFYKTEIKNRVEFESALYPNW
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                                                                                                                                                                                                                                                                    39;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
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                                                                                                                            ETDKIPVALGLKEKNLYLSCGMKDGKPTLQLETVD-PNTYPKRKME 157
                                                                                                                                                                                                                                                                                                15.1%;
33.3%;
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28.7%;
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                                                                                                                                                                                                                                                                                                Score 124.5;
Pred. No. 2.4
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Pred. No. 1.1e-05;
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                                                                                                                                                                                                                                                                48;
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A; Reference number: I51852
A; Accession: I51852
A; Accession: I51852
A; Status: translated from GB/EMBL/DDBJ
A; Molecule type: mRNA
A; Molecule type: mRNA
A; Molecule type: mRNA
A; Residues: I-5, 'K', 7-19, 'H', 21-110, 'Q', 112-176, 'A', 178-213, 'P', 215-269 <
A; Cross-references: GB:M54933; NID:9186287; PIDN:AAA59136.1; PID:9186288
R; Nishida, T.; Nishino, N.; Takano, M.; Kawai, K.; Bando, K.; Masul, Y.;
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A; Title: GDNA cloning of IL-1 alpha and IL-1 beta from mRNA of U937 cell
A; Reference number: I52217; MUID:87156769
A; Accession: I65200
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C;Species: Homo sapiens (man)
C;Date: 28-reb-1986 #sequence_revision 15-May-1998 #text_change 15-Sep-2000
C;Accession: A25542; A29019; A94023; A93361; I51852; I65200; I38132; B27616;
R;Clark, B.D.; Collins, K.L.; Gandy, M.S.; Webb, A.C.; Auron, P.E.
Nucleic Acids Res. 14, 7897-7914, 1986
Nucleic Acids Res. 14, 7897-7914, in prointerleukin 1 beta: possible evolution
A;Cross-references: EMBL:X56087; NID:g35662; PIDN:CAA39567.1; PR:Zsebo, K.M.; Wypych, J.; Yuschenkoff, V.N.; Lu, H.; Hunt, P.; Blood 71, 962-968, 1988
                                                                                                                                                                                                                                                                                                                                            A;Cross-references: GB:M15330; NID:g186283; PIDN:AAA59135.1; R;Kotenko, S.V.; Bulenkov, M.T.; Veiko, V.P.; Epishin, S.M.; ii, S.A.; Vinetskii, Y.P.
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A;Residues: 1-269 <NIS>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        A; Molecule type: mRNA
A; Residues: 1-5, K, 7-269 < AUR>
A; Residues: 1-5, K, 7-269 < AUR>
A; Cross-references: GB: K02770; NID: g186268; PIDN: AAA36106.1; PID: g307043
R; March, C.J.; Mosley, B.; Larsen, A.; Cerretti, D.P.; Braedt, G.; Price,
Nature 315, 641-647, 1985
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A; Title: Human interleukin-1 beta gene.
A; Reference number: A29019; MUID:87248099
A; Accession: A29019
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A;Note: the mRNA sequence had codon AAG for
R;Bensi, G.; Raugei, G.; Palla, E.; Carinci,
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A;Title: [Cloning of the cDNA coding for l
A;Reference number: 138131; MUID:90249285
A;Accession: 138132
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A; Residues: 1-269 <MAR>
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A; Residues: 1-269 <BEN>
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A; Residues: 1-5, 'K', 7-269 <CLA>
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N;Alternate names: hematopoietin-1; IL-1 be
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Lomakin, I.B.;
                                                                 PID: g35663
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F;10/Binding site: myristate (Lys) (covalent) (partial) #status experimental
F;103/Binding site: carbohydrate (Asn) (covalent) #status absent
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A;Map position: 2q13-2q21
A;Introns: 16/2; 33/3; 101/1; 156/1; 199/3
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J. Biol. Chem. 266, 7081-7086, 1991
A;Title: Structure-function mapping of interleukin 1 precursors.
A;Reference number: A39774; MUID:91201363
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A;Title: The 31-kDa precursor of interleukin lalpha is myristoylated on specific lysines A;Reference number: A48293; MUID:93348250
A;Contents: annotation; myristylation of lysines A;Contents: annotation; myristylation of lysines R;Nanduri, V.B.; Hulmes, J.D.; Pan, Y.C.E.; Killan, P.L.; Stern, A.S. Biochim. Biophys. Acta 1118, 25-35, 1991
                                                                        RESULT
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interleukin-1 beta precursor
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Pred. No. 4.6e-05;
3; Mismatches 61
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A; Reference number: A24719; MUID:87117546
A; Accession: A24719
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C:Species: Oryctolagus cuniculus (domestic rabbit) (C:Date: 25-May-1989 #sequence_revision 22-Nov-1996 #text_change C:Accession: A27714; A30584; JUON82; A32166 R:MOT1, S:; Goto, F:; Goto, K.; Ohkawara, S:; Maeda, S:; Shimada Blochem. Blophys. Res. Commun. 150, 1237-1243, 1988 A;Title: Cloning and sequence analysis of a cDNA for lymphocyte A;Reference number: A27714; MUID:88134238
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A.Rosidues: 1-268 <YOU>
C.Comment: This protein lacks a conventional signal sequence for protein export.
ved form of interleukin-lbeta, unlike interleukin 1-alpha, is inactive.
C.Comment: Interleukin-lbeta precursor is less heavily myristoylated than interle
                                                                                                                                                                                                                C;Species: Mus musculus (house mouse)
C;Date: 26-Uul-1996 #sequence_revision 22-Nov-1996 #text_change 22-Jun C;Date: 26-Wi-1996; A24719; S13029
R;Gray, P.W.; Glaister, D.; Chen, E.; Goeddel, D.V.; Pennica, D. J. Immunol. 137, 3644-3648, 1986
A;Title: Two interleukin 1 genes in the mouse: Cloning and expression A;Reference number: IS5969; MUID:87058957
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A;Title: Rabbit II-1. Cloning, expression, biologic properties, A;Reference number: A30584; MUID:89176242
A;Accession: A30584
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Protein Eng. 2, 545-551, 1989
A;Title: Cloning of rabbit interleukin-1 beta:
A;Reference number: A94230; MUID:89315718
A;Accession: JU0082
                       R; Telford, J.L.; Macchia, G.; Massone, Nucleic Acids Res. 14, 9955-9963, 1986
                                                                            A; Cross-references: GB:M15131; NID:g198293; PIDN:AAA39276.1; PID:g309398
                                                                                                               A; Residues: 1-269 < RES>
                                                                                                                                      A; Molecule type: mRNA
                                                                                                                                                                    A; Status: preliminary;
                                                                                                                                                                                           A; Accession: I55969
                                                                                                                                                                                                                                                                                                                                                                                                              interleukin-1 beta precursor - mouse
N;Alternate names: hematopoietin-1; IL-1 beta
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           C;Keywords = Cytokine; immunoregulation; inflammation; lymphokine; macrophage; mitogen F:117-268/Product: interleukin-1 beta #status predicted <ILB>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        A; Residues: 1-268 <CAN>
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A; Residues: 1-268 < MOR>
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A; Title: The murine interleukin 1-beta gene: structure and evolution
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   A;Cross-references: GB:M26295; NID:g516632;
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    133 LVLSGTY-----ELKALHLNAENLNQQVVFSMSFVQGEESN------DKIPVALG 176
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   VQGGSQCLSCGV-GQEPTLTLEPVNIMELYLGAKESKSFTFYRRDMGLTSSFESAAYPGW 119
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          44;
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Pred. No. 9.3e-05;
1; Mismatches 63
                                                    Α.,
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                                                 Carinci,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PIDN: AAA31373.1;
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C;Species: Cyprinus carpio (common carp)
C;Date: 22-Oct-1999 #sequence_revision 22-Oct-1999 #text_change 22-Oct-1999
C;Date: 22-Oct-1999 #sequence_revision 22-Oct-1999 #text_change 22-Oct-1999
C;Accession: T30517
R;Nakao, M.; Obo, R.; Mutsuro, J.; Fujiki, K.; Nonaka, M.; Yano, T.
R;Nakao, M.; Obo, R.; Mutsuro, J.; Fujiki, K.; Nonaka, M.; Yano, T.
R;Description: Sequence diversity of the third complement component (C3) from a bony fis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            A;Gene: IL-1-beta
C;Superfamily: interleukin-1
C;Reywords: cytokine; immunoregulation; inflammation; lymphokine; macrophage; mitogen
F;118-269/Product: interleukin-1 beta #status experimental <IL1>
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A;Residues: 118-269 <CAND
C;Comment: This protein lacks a conventional signal sequence for protein export. Cleavag
ved form of interleukin-lbeta, unlike interleukin 1-alpha, is inactive.
C;Comment: Interleukin-lbeta precursor is less heavily myristoylated than interleukin-la
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A;Residues: 1-269 <TEL>
A;Cross-references: GB:X04964; NID:952666; PIDN:CAA28637.1; PID:952667
A;Cross-references: GB:X04964; NID:952666; PIDN:CAA28637.1; PID:952667
A;Campy, G.O.; Wilder, C.L.; Merenda, J.M.; McColl, A.S.; Geoghegan, K.F.; Otterness, I.
FEBS Lett. 278, 98-102, 1991
A;Title: Reduction of biological activity of murine recombinant interleukin-1beta by sel
A;Reference number: S13029; MUID:91130610
A;Accession: S13029; MUID:91130610
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A;Molecule type: mRNA
A;Residues: 1-1230 <NAK>
A;Cross-references: EMBL;AB016215; NID:d1253045; PID:d1037610; PIDN:BAA36623.1
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A; Accession: T30517
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Best Local S
Matches 48
                                                                                                                                                                                                                                                                                                                                  Ouery Match 9.8%; Score 80.5; DB 2; Best Local Similarity 27.3%; Pred. No. 7.3; Matches 35; Conservative 17; Mismatches 53.
                                                  118 GWFLCTVP 125
                                                                                                                                                                                                                          673
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                                                                                                        733 ASLHESIRKAVSFLEGRLPQLINPYAVAMMSYAMANENK----LNKDILMKHSSQHEAGR 788
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                                                                                                                                                                                                                                                                            14 SALKVLYLHNNQLLAGGLHAGKVIKGEEISVVPNRWLDASLSP-VILGVQGGSQCLSCGV 72
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SW---TVP 793
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                                                                                                                                                              GQEPTLTLEPVNIMELYLGAKESKSFTFYRRDMGLTSSFESAAYP 117
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                                                                                                                                                                                                                                                                                                                                     53;
                                                                                                                                                                                                                                                                                                                                                                                        Length 1230;
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Search completed: June 25, 2001, 14:05:59
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Maximum Match 1008
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           Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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MEDLINE-90354444; PubMed-2143761;
Blenkowski M.J., Eessalu T.E., Berger A.E., Truesd Shelly J.A., Laborde A.L., Zurcher-Weely H.A., Rea Heinrikson R.L., Chosay J.G., Tracey D.E.;
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MEDLINE=90136920;
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                                                                                                                                                                                                                                                                                                                                 Biol. Chem.
                                         FUNCTION: IL-1RA INHIBITS THE ACTIVITY OF IL-1 BY BINDING TO RECEPTOR. IL-1RA HAS NO IL-1 LIKE ACTIVITY.
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MEICRGLRSHLITLLLFLFHS
INTRACELLULAR ISOFORM)
D1690776A7394057 CRC6
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                                                                                                            153
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Created)

STANDARD; 22,

PRT;

178 8

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MEDLINE-91271363; ruproc. Elsenberg S.P., Brewer M.T., Verderber E., nearly Brandhuber B.J., Thompson R.C.;
"Interleukin 1 receptor antagonist is a member of the gene family: evolution of a cytokine control mechanism gene family: evolution of a cytokine control mechanism gene family:
         Pfam; PF00340; interleukin-1; 1. PRINTS; PR00264; INTERLEUKIN1. PROSITE; PS00253; INTERLEUKIN_1;
                                                                                                 EMBL;
EMBL;
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PIR; I
                                                                                                                                                                                                                           modified and this statement is not removed. Usage by and frentities requires a license agreement (See http://www.isb-sibor send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                               This SWISS-PROT entry is copyright. It is produced between the Swiss Institute of Bioinformatics and the European Bioinformatics Institute. There are r
                                                                                                                                                                                                                                                                                                                                                                                                                              SEQUENCE OF 23-178 FROM N.A.
MEDLINE=92037824; PubMed=1834470;
Shuck M.E., Eessalu T.E., Tracey D.E.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE FROM N.A.

MEDLINE=91316273; PubMed=1830498;

Matsushime H., Roussel M.F., Matsushima K., Hishinuma A.,

"Cloning and expression of murine interleukin-1 receptor
in macrophages stimulated by colony-stimulating factor 1.

Blood 78:616-623(1991).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE FROM N.A.

MEDLLINE=91250712; PubMed=1828262;

MEDLINE=91250712; PubMed=1828262;

Zahedi K., Seldin M.F., Rits M., Ezekowitz R.A

"Mouse IL-1 receptor antagonist protein. Molec

gene mapping, and expression of mRNA in vitro

J. Immunol. 146:4228-4233(1991).
                                                                                                                                                                                                                                                                                                                                                                                  "Cloning, heterologous expression and characterization of murine interleukin 1 receptor antagonist protein.";
Eur. J. Immunol. 21:2775-2780(1991).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE OF 7-178 FROM N.A.
MEDLINE=91271363; PubMed=1828896;
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Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              01-MAY-1992
01-NOV-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MEDLINE=94271931; PubMed=8003626;
Zahedi K.A., Uhlar C.M., Rits M.,
"The mouse interleukin 1 receptor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NCBI_TaxID=10090;
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                                                      nterPro; IPR000975;
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                                                                                                ; B40956;
; A44610;
                                                                                                                                                                                                                                                                                                                                          FUNCTION: IL-1RA INHIBITS THE ACTIVITY OF IL-1 RECEPTOR. IL-1RA HAS NO IL-1 LIKE ACTIVITY. SIMILARITY: BELONGS TO THE IL-1 FAMILY.
                                                                                                                                                                                                                                                                                                                                                                  FUNCTION:
                                                                                                                         L; M74294; AAA39309.1; -.
L; M64004; AAA39277.1; -.
L; L32838; AAA20576.1; -.
L; M57525; AAA39278.1; -.
L; M63100; AAA39310.1; -.
L; S64082; AAB20265.1; -.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         regulation in vitro.";
okine 6:1-9(1994).
                                                                #GI:96547; Illrn.
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Rodentia;
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Last annotation updat
R ANTAGONIST PROTEIN P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              tion update)
PROTEIN PRECURSOR (IL-1RA) (IL-1RN)
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thi; Muridae;
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; Murinae; Mus.
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Matches 71
    DISULFID CARBOHYD SEQUENCE
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15-DEC-1998
15-DEC-1998
15-JUL-1999
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CARBOHYD
SEQUENCE
                                                                                                                                                  the European Bioinformatics Institute. The use by non-profit institutions as long modified and this statement is not removed.
                                                                                                                                                                                                                      "Characterization of IRAP in morphine treated pig.";
Submitted (MAY-1996) to the EMBL/GenBank/DDBJ databases
-i-FUNCTION: IL-1RA INHIBITS THE ACTIVITY OF IL-1 BY B:
RECEPTOR. IL-1RA HAS NO IL-1 LIKE ACTIVITY.
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CHAIN
                                                                                                                                entities requires a license agreement (Su or send an email to license@isb-sib.ch).
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                                                                                                                                                                                                                                                                                                              Sus scrofa (Pig).
Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                     Glycoprotein;
SIGNAL
                                                                                            Pfam;
                                                                                                                EMBL; L38849; AAA99424.1;
                                                                                                                                                                                                                                                                         STRAIN=CROSSBREED;
                                                                         PROSITE; PS00253;
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                                                                                                      [nterPro;
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                                                                                                                                                           SWISS-PROT entry is copyright. It is produced through a cen the Swiss Institute of Bioinformatics and the EMBL European Bioinformatics Institute. There are no restrict by non-profit institutions as long as its content is
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                                                                                            PF00340;
                                                                               PR00264; INTERLEUKIN1.
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37, Last sequence update)
38, Last annotation update)
;EPTOR ANTAGONIST PROTEIN PRECURSOR (IL-1RA) (IL-1RN)
                                                                       INTERLEUKIN_1;
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177
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                       PROTEIN.
BY SIMILARITY.
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                                           INTERLEUKIN-1
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No. 6.2
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                                           RECEPTOR ANTAGONIST
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. .; (POTENTIAL)
CRC64;
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                                                                                                                                                                        restrictions
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Query Match Best Local Similarity

37.1%; 48.6%;

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305.5; DB 1; No. 1.3e-24;

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CARBOHYD
SEQUENCE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              077482;
                                                                                                                                                             GIYCOProtein; Signal.
SIGNAL 1 2
                                                                                                                                                                                 PROSITE;
                                                                                                                                                                                                     Pfam; PF00340;
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InterPro; IPR000975; -.
                                                                                                                                                                                                                                                                                                                                                            "Enzymatic amplification and expression of bovine receptor antagonist cDNA.";
                                                                                                                                                                                                                                                                                                                                                                                          Kirisawa R.,
                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE FROM N.A. MEDLINE-98305607;
                                                                                                                                                                                                                                                                                                                                                                                                                                  NCBI_TaxID=9913
                                                                                                                                                                                                                                                                                                                                                                                                                                            Bovidae; Bovinae; Bos.
                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mammalia; Eutheria;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    IL1RN
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15-JUL-1999 (Rel. 38, Last annotation update)
INTERLEUKIN-1 RECEPTOR ANTAGONIST PROTEIN PRECURSOR (IL-1RA) (IL-1RN)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          15-DEC-1998
15-DEC-1998
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                                                                                                                                                                                                                                                                                                                      . Immunol. Immunopathol. 62:197-208(1998).
FUNCTION: IL-1RA INHIBITS THE ACTIVITY OF IL-1 BY BINDING TO RECEPTOR. IL-1RAHAS NO IL-1 LIKE ACTIVITY.
SIMILARITY: BELONGS TO THE IL-1 FAMILY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        taurus (Bovine).
CLSC-GVGQEPTLTLEPVNIMELYLGAKESKSFTFYRRDMGLTSSFESAAYPGWFLCTVP 125
                                FRMKDSALKVLYLHNNQLLAGGLHAGKVIKGEEISVVPNRWLDASLSP--VILGVQGGSQ 66
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                   FRIWDVNQKIFYLRNNQLVAGYLQGPNTKLEEKIDVVP-----IEPHTMFLGIHGGKL 88
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                                                                                                                                                                                          PR00264; INTERLEUKIN1
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                                                                   Similarity
                                                                                                                                                                                PS00253; INTERLEUKIN_1; 1.
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174
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174
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                                                                 35.9%;
47.3%;
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                                                                                                            MW.
                                                           14;
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                                                                   Score 295.5;
Pred. No. 1.
                                                                                                        BY SIMILARITY.
N-LINKED (GLCNAC. . .) (POTENTIAL); 1E56E7F224FF851F CRC64;
                                                                                                                                          PROTEIN
                                                                                                                                                  INTERLEUKIN-1 RECEPTOR ANTAGONIST
                                                                                                                                                             BY SIMILARITY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PRT;
                                                           Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mismatches
                                                                                                                                                                                                                                                                                                                                                                                           Hagiwara K.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       174 AA
                                                          .3e-23;
les 51;
                                                                             DB 1;
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                                                                            Length
                                                          Indels
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                                                                               174;
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SEQUENCE
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154
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                  128
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01-MAY-1992
01-NOV-1995
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Eukaryota; Métazoa; Chordata;
Mammalia; Eutheria; Rodentia;
NCBI_TaxID=10116;
                                                                                                                                                                                                                                                                                                                                                                                          Glycoprotein; Signal. SIGNAL 1 2
                                                                                                                                                                                                                                                                                                                                                                                                                          PRINTS; PR00264; INTERLEUKIN1.
PROSITE; PS00253; INTERLEUKIN_1; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             EMBL; M63101; AAA41434.1;
PIR; C40956; C40956.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Brandhuber B.J., Thompson R.C.;
"Interleukin 1 receptor antagonist is a member of the interleukin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Eisenberg S.P., Brewer M.T.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Rattus norvegicus (Rat).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        INTERLEUKIN-1 RECEPTOR ANTAGONIST PROTEIN PRECURSOR (IL-1RA) (IL-1RN)
                                                                                                                                                                                                                                                                                                                                                                                                                                                             InterPro; IPR000975;
Pfam; PF00340; inter
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                HSSP; P18510;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SIMILARITY: BELONGS TO THE IL-1 FAMILY.
DHPVSLTNTPKE - - - PCTVTKFYFQE
                                  DOPVRLTQLPENGGWNAPITDFYFQQ 153
                                                                                                                                                                         FRMKDSALKVLYLHNNQLLAGGLHAGKVIKGEEISVVPNRWLDASLSPVILGVQGGSQCL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 EADQPVGLTNMPTEA --- LKVTKFYFQQ 173
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    EADQPVRLTQLPENGGWNAPITDFYFQQ 153
                                                                                     SC-GVGQEPTLTLEPVNIMELYLGAKESKSFTFYRRDMGLTSSFESAAYPGWFLCTVPEA
                                                                                                                                        FRIWDTNQKTFYLRNNQLIAGYLQGPNTKLEEKIDMVP----IDFRNVFLGIHGGKLCL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CLACVKSGDEIKLKLEAVNITDLNQNREQDKRFAFIRFDNGPTTSFESAACPGWFLCTSL 148
                                                                   SCVKSGDDTKLQLEEVNITDLNKNKEEDKRFTFIRSETGPTTSFESLACPGWFLCTTLEA
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45.9%;
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Last annotation update)
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                                                                                                                                                                                                                                                                                                      MW.
                                                                                                                                                                                                              14;
                                                                                                                                                                                                        Score 292.5; DB 1;
Pred. No. 2.8e-23;
                                                                                                                                                                                                                                                                                                BY SIMILARITY.

N-LINKED (GLCNAC. . .) (P

F3A5754FB6C51B03 CRC64;
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                                                                                                                                                                                                                                                                                                                                                                                          BY SIMILARITY.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Craniata; Vertebrata; I
Sciurognathi; Muridae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PRT;
 176
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (See http://www.isb-sib.ch/announce/
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 There are no restrictions ng as its content is in
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; Murinae; Rat
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                                    Query Match
Best Local
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                                                                                    DISULFID
CARBOHYD
SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  P26890;
01-AUG-1992 (Rel. 23, Created)
01-AUG-1992 (Rel. 23, Last sequence update)
15-DEC-1998 (Rel. 37, Last annotation update)
INTERLEUKIN-1 RECEPTOR ANTAGONIST PROTEIN PRE
                                                                                                                                                                                                                                   EMBL; $68977; AAB30093.1; -.
EMBL; M57526; AAA31374.1; -.
EMBL; D21832; BAA04860.1; -.
EMBL; A54377; A54377.
                                                                                                                                                                                                                                                                                                   entities requires a license agreement (S or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                           use by non-profit institutions as long modified and this statement is not removed.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                Goto F., Goto K., Miyata T., Ohkawara S., Takao T., Mori S., Furukawa S., Maeda T., Iwanaga S., Shimonishi Y., Yoshinaga M., "Interleukin-1 receptor antagonist in inflammatory exudate cells rabbits. Production, purification and determination of primary
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MEDLINE=93052512; PubMed=1427977; Ohka
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE FROM N.A. Hamada H., Mulligan | Submitted (XXX-1992)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Cominelli F., Bortolami M., Pizarro T
Brewer M.T., Eisenberg S.P., Ng R.K.;
"Rabbit interleukin-1 receptor antago
                                                                                                                                                 Glycoprotein;
SIGNAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE FROM N.A. MEDLINE=94165101; PubMed=7509813;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Oryctolagus cuniculus (Rabbit).
Eukaryota; Metazoa; Chordáta; Craniata;
                                                                                                                                                                                                                                                                                                                                                  the European Bioinformatics Institute.
                                                                                                                                                                                                                                                                                                                                                                                                                                                        structure."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               functional characterization, and regulation during intestinal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      IL1RN OR IL1RA.
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                                                                                                                                                                                                                                                                                                                                                                                                  FUNCTION: IL-1RA INHIBITS THE ACTIVITY OF I RECEPTOR. IL-1RA HAS NO IL-1 LIKE ACTIVITY. SIMILARITY: BELONGS TO THE IL-1 FAMILY.
FRMKDSALKVLYLHNNQLLAGGLHAGKVIKGEEISVVPNRWLDASLSP--VILGVQGGSQ 66
                                                                                                                                                                                                   PF00340;
                                                                                                                                                                                                                            P18510; 1ILR.
                                     Similarity
                                                                                                                                                                         PR00264; INTERLEUKIN1.
; PS00253; INTERLEUKIN_1; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Eutheria;
                                                                                                                                                                                                             IPR000975; -
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109
177
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                          Conservative
                                                                                                                                      ; Signal.
1 25
26 177
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AAA31374.1; -.
                                                                                                                                                                                                interleukin-1; 1.
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109
20214 MW;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             receptor antagonist. Cloning,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        the EMBL/GenBank/DDBJ databases
                          17;
                                                                                PROTEIN.
BY SIMILARITY.
N-LINKED (GLCNAC. .
W; F5BC087F097FFEAF C
                        Score 286.5;
Pred. No. 1.2e
7; Mismatches
                                                                                                                                                 BY SIMILARITY.
                                                                                                                                      INTERLEUKIN-1 RECEPTOR ANTAGONIST
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Vertebrata; Eu
ae; Oryctolagus
                                   ; DB 1
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                                                                                     CRC64;
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                        Indels
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                                                                                                (POTENTIAL)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             expression,
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                        13;
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IL1X_HO
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Query Match
Best Local Similarity
Matches 65; Conser
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018999; 077745;

15-DEC-1998 (Rel. 3

15-DEC-1998 (Rel. 3

15-JUL-1999 (Rel. 3
                                                                                                                                                DISULFID CARBOHYD CONFLICT
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SEQUENCE FROM N.A.
MEDLINE-97366446; PubMed-9223227;
Kato H., Ohashi T., Matsushiro H.,
Haseqawa A.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           _HORSE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            This SWISS-PROT entry is copyright. It is produced through between the Swiss Institute of Bioliformatics and the Entropean Bioinformatics Institute. There are no restruct by non-profit institutions as long as its content use by non-profit institutions as long as its content modified and this statement is not removed. Usage by and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   "Cloning of equine interleukin-1 receptor antagonist and determination of its full-length cDNA sequence."; Am. J. Vet. Res. 59:712-716(1998).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Eukaryota; Metazoa;
Mammalia; Eutheria;
NCBI_TaxID=9796;
                                                                                                                                                                                                                                                           CHAIN
                                                                                                                                                                                                                                                                                      Glycoprotein; SIGNAL
                                                                                                                                                                                                                                                                                                                                                                                                                                    EMBL; D83714; BAA22529.1; -. EMBL; U92482; AAC39257.1; -.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               entities requires a license agreement (S or send an email to license@isb-sib.ch).
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15-DEC-1998 (Rel. 37, Last sequence update)
15-DEC-1999 (Rel. 38, Last annotation update)
INTERLEUKIN-1 RECEPTOR ANTAGONIST PROTEIN PRECURSOR (IL-1RA) (IL-1RN)
                                                                                                                             SEQUENCE
                                                                                                                                                                                                                                                                                                                                   PRINTS; PR00264; INTERLEUKIN1.
PROSITE; PS00253; INTERLEUKIN_1; 1.
                                                                                                                                                                                                                                                                                                                                                                                      Ptam;
                                                                                                                                                                                                                                                                                                                                                                                                              InterPro; IPR000975;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        -!- SIMILARITY: BELONGS TO THE IL-1 FAMILY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      MEDLINE=98285942; PubMed=9622739;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Tsujimoto H., Hasegawa A.;
"Molecular cloning and functional
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Equus caballus (Horse).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           receptor antagonist."
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  OF VEC. RES. 39:/12-7/10(1998).
EUNCTION: IL-IRA INHIBITS THE ACTIVITY OF IL-1
RECEPTOR. IL-IRA HAS NO IL-1 LIKE ACTIVITY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Immunol.
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109
19
                                                                                                                                                                                                                                                                                                           Signal
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                                                                                                                             AA;
                                                                                                                                                                                                                                                                                                                                                                                    interleukin-1;
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                        33.2%;
                                                                                                                             20459
                                                                                                                           MW;
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F -> L (IN REF. 2).
; 1ABC377F1F1CF80B
                     Score 273.5;
Pred. No. 2.6
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BY SIMILARITY
                                                                                                                                                                                                                                                         BY SIMILARITY.
INTERLEUKIN-1 RECEPTOR ANTAGONIST
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                                             Length 177;
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                                                                                      EMBL; X54796; CAA38566.1; -
EMBL; X56972; CAA40293.1; -
PIR; S13092; S13092.
PIR; S13810; S13810.
PIR; S23010; S23010.
HSSP; P01584; 411B.
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P21621;
01-MAY-1991 (Rel. 18, Created)
01-MAR-1992 (Rel. 21, Last sequence updat
30-MAY-2000 (Rel. 39, Last annotation upd
INTERLEUKIN-1 BETA PRECURSOR (IL-1 BETA).
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[2]
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"Nucleotide sequence of ovine interleukin-1 beta.";
Nucleic Acids Res. 18:7165-7165(1990).
-i- FUNCTION: PRODUCED BY ACTIVATED MACROPHAGES, IL-1 STIMULATES
THYMOCYTE PROLIFERATION BY INDUCING IL-2 RELEASE, B-CELL
MATURATION & PROLIFERATION BY INDUCING IL-1
MATURATION & PROLIFERATION IN THE INFLAMMATORY RESPONSE, BEING
IL-1 PROTEINS ARE INVOLVED IN THE INFLAMMATORY RESPONSE, BEING
IDENTIFIED AS ENDOGENOUS PYROGENS, AND ARE REPORTED TO STIMULATE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Cetartiodactyla; Ruminantia; Pecora; Bovoidea; Bovidae; Caprinae; Ovis.
NCBI_TaxID=9940;
PRINTS;
                                 InterPro; IPR000975; -.
InterPro; IPR002348; -.
Pfam; PF00340; interleukin-1;
                                                                                                                                                                                                                         entities requires a license agreement (See or send an email to license@isb-sib.ch).
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Seow H.F., Rothel J.S., David M.J., Wood P.R.;
"Nucleotide sequence of ovine macrophage interleukin-1 beta cDNA.";
DNA Seq. 1:423-426(1991).
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                                                                                                                                                                                                                                                        European Bioinformatics Institute. The by non-profit institutions as long ified and this statement is not removed.
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MISCELLANGOUS: THE LACK OF A SPECIFIC TUDROPHOBIC SEGMENT IN THE PRECURSOR SEQUENCE SUGGESTS THAT IL-1 IS RELEASED BY DAMAGED CELLS OR IS SECRETED BY A MECHANISM DIFFERING FROM THAT USED FOR OTHER SECRETED BY A MECHANISM DIFFERING FROM THAT USED FOR OTHER
                                                                                                                                                                                                                                                                                                                                                                          SECRETORY PROTEINS.
SIMILARITY: BELONGS TO THE IL-1 FAMILY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SUBUNIT: MONOMER.
                                                                                                                                                                                                                                                                              SWISS-PROT entry is copyright. It is produced through a collaboration een the Swiss Institute of Bioinformatics and the EMBL outstation European Bioinformatics Institute. There are no restrictions on in by non-profit institutions as long as its content is in no war by non-profit institutions as long as its content is in no war by non-profit institutions.
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                   PR00262;
                   IL1HBGF
INTERLEUKIN1
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Last annotation updat
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Matches 43
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                                                           modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
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01-OCT-1996 (Rel. 34, Last sequence updat
30-MAY-2000 (Rel. 39, Last annotation upd
INTERLEUKIN-1 BETA PRECURSOR (IL-1 BETA).
                        EMBL; U20500; AAA62234.1; HSSP; P01584; 1HIB.
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                                                                                                                                                                                                                                                                                                                                                                          Submitted (MAR-1995)
                                                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE
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Cervidae; Cervinae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Eukaryota;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Cervus elaphus (Red
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       P51745;
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           nterPro;
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                                                                                                                                                                           DOMAIN: THE SIMILARITY AMONG THE IL-1 PRECURSORS SUGGESTS THAT THE AMINO ENDS OF THESE PROTEINS SERVE SOME AS YET UNDEFINED FUNCTION. MISCELLANDOUS: THE LACK OF A SPECIFIC HYDROPHOBIC SEGMENT IN THE PRECURSOR SEQUENCE SUGGESTS THAT IL-1 IS RELEASED BY DAMAGED CELLS OR IS SECRETED BY A MECHANISM DIFFERING FROM THAT USED FOR OTHER SECRETORY PROTEINS.
                                                                                                                                                                                                                                                                                                   mitted (MAR-1995) to the EMBL/GenBank/DDBJ databases. EUNCTION: PRODUCED BY ACTIVATED MACROPHAGES, IL-1 STIMULATES THYMOCYTE PROLIFERATION BY INDUCING IL-2 RELEASE, B-CELL MATURATION & PROLIFERATION, & FIBROBLAST GROWTH FACTOR ACTIVITY. IL-1 PROTEINS ARE INVOLVED IN THE INFLAMMATORY RESPONSE, BEING IDENTIFIED AS ENDOGENOUS PYROGENS, AND ARE REPORTED TO STIMULATE
                                                                                                                        European
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                                                                                                                                   SWISS-PROT entry is copyright. It is produced through a collaboration -
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43; Conserv
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                                                                                                        pean Bioinformatics Institute. The non-profit institutions as long
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31.9%;
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Q -> K (IN I
V -> A (IN I
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Y -> C (IN RE
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(IN REF. 2).
(IN REF. 2).
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                                                                                                                                                                                                                                                                                          COLLAGENASE FROM
                                                                                                                                                                                                                                                                                                                                                                                                                                                          Ruminantia; Pecora; Cervoidea;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    266
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                                                                                                      There are no restrictions ng as its content is in
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RESULT 10
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DT 01-MAR
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DE III.B.
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OC ELKAZY
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PRINTS; PR00262; ILLHBGE.
PRINTS; PR00264; INTERLEUKIN.1; 1.
PROSITE; PS00253; INTERLEUKIN.1; 1.
Cytokine; Macrophage; Mitogen; Inflammatory response;
PROPEP 1 113 BY SIMILARITY.
PROPEP 1 114 266 INTERLEUKIN-1 BETA.
PROPER 114 266 AF40B4E6F0D9F060 CRC64
                                This SWI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Eukaryota; Metazoa;
Mammalia; Eutheria;
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01-MAR-1989
                                                                                                                                                                                                                                                                                                                                                                                                                      "Cloning, sequence and expression of bovine interleukin 1 beta complementary DNAs."; Mol. Immunol. 25:429-437(1988).
-i- FUNCTION: PRODUCED BY ACTIVATED MACROPHA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        01-MAR-1989
30-MAY-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gillis S., Cerretti D.P.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MEDLINE=88318652; PubMed=3261832; Maliszewski C.R., Baker P.E., Sch
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Nucleic Acids
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 01-MAR-1989 (Rel. 10, Created)
01-MAR-1989 (Rel. 10, Last seq
30-MAY-2000 (Rel. 39, Last ann
INTERLEUKIN-1 BETA PRECURSOR (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Bovidae; Bovinae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       IL1B_BOVIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             249
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                                                                                                                                              DOMAIN: THE SIMILARITY AMONG THE IL-1 PRECURSORS SUGGESTS THAT THE AMINO ENDS OF THESE PROTEINS SERVE SOME AS YET UNDEFINED FUNCTION.
MISCELLANGOUS: THE LACK OF A SPECIFIC HYDROPHOBIC SEGMENT IN THE PRECURSOR SEQUENCE SUGGESTS THAT IL-1 IS RELEASED BY DAMAGED CELLS.
OR IS SECRETED BY A MECHANISM DIFFERING FROM THAT USED FOR OTHER
                                                                                                    SECRETORY PROTEINS.
SIMILARITY: BELONGS
                                                                                                                                                                                                                                                                                        FUNCTION: PRODUCED BY ACTIVATED MACROPHAGES, IL-1 STIMULATES THYMOCYTE PROLIFERATION BY INDUCING IL-2 RELEASE, B-CELL MATURATION & POLIFERATION, & FIBROBLAST GROWTH FACTOR ACTIVITY. BROTELING ARE INVOLVED IN THE INFLAMMATORY RESPONSE, BEING IDENTIFIED AS ENDOGENOUS PYROGENS, AND ARE REPORTED TO STIMULATE THE RELEASE OF PROSTAGLANDIN AND COLLAGENASE FROM SYNOVIAL CELLS.
                                                                                                                                                                                                                                                                         THE RELEASE OF PROSTAGLANDIN AND SUBUNIT: MONOMER.
                         SWISS-PROT entry is copyright. It is produced through a centre the Swiss Institute of Bioinformatics and the EMBL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             HF--RGGQD--ITDF 259
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       EPTLTLEPVNIMELYLGAKESKSFTFYRRDMGLTSSFESAAYPGWFLCTVPEADQPVRLT 134
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          Swiss Institut
Bioinformatics
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Cetartiodactyla;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            17.1%;
31.9%;
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Last annotation update)
                                                                                                         TO
                                                                                                      THE IL-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     22;
       Institute.
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Pred. No. 2.6e
22; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2 140.5;
                                                                                                      FAMILY.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Davis
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       no
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                                                  a collaboration
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   15;
     outstation -
cions on its
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IL1B_HORSE
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Best Local
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PRINTS; PRO0264; INTERLEUKIN1.; 1.
PROSITE; PS00253; INTERLEUKIN_1; 1.
CYCOKINE; Macrophage; Mitogen; Inflam
PRODEP 1 113
CHAIN 114 266
CONFLICT 252 252
SEQUENCE 266 AA; 30774 MW; 9D1EF8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    EMBL; M35589; AAA30585.1; -.
EMBL; X12498; CAA31018.1; -.
EMBL; M37211; AAA30584.1; -.
PIR; JL0010; ICBOLB.
PIR; S01380; S01380.
HSSP; P01584; 1HIB.
                                                                                                                                                                                                                                                      Howard R.D., McIlwraith C.W., Trotter G
"Cloning of equine interleukin-1 alpha
and determination of their full-length
Am. J. Vet. Res. 59:704-711(1998).
                                                                                                                                                                                                               Eukaryota; Metazoa;
Mammalia; Eutheria;
                  SEQUENCE FROM N.A. (SHORT FORM).
MEDLINE=97080493; PubMed=8921838;
                                                                                                                                                                                                                                    Equus caballus (Horse).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             entities requires a license agreement (Some send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  use by non-profit institutions as long modified and this statement is not removed
           Kato H.,
                                                                                            SEQUENCE FROM N.A.
MEDLINE=98285941; PubMed=9622738;
                                                                                                                                                Tsujimoto H., Hasegawa A.;
                                                                                                                                                             Kato
                                                                                                                                                                     SEQUENCE FROM N.A.
MEDLINE=96131982; PubMed=8578682;
                                                                                                                                                                                                     NCBI_TaxID=9796;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Pfam; PF00340;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                InterPro; IPR000975; -.
InterPro; IPR002348; -.
                                                                                                                                     Molecular cloning
                                                                                                                                                                                                                                                                                                                                                                  248
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                                                                                                                                                             H., Ohashi T.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         LKYLYLHNNQLLAGGLHAGKVIKGEEISVVPNRWLDASLSPVILGVQGGSQCLSC-GVGQ
                                                                                                                              Immunol.
                                                                                                                                                                                                                                                                                                                                                                                      QLPENGGWNA - - PITDF
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                                                                                                                                                                                                                                                                                                                                                                                                                              EPTLTLEPVNIMELYLGAKESKSFTFYRRDMGLTSSFESAAYPGWFLCTVPEADQPVRLT 134
                                                                                                                                                                                                                                                                                                                                                                                                                                                    LKALHLLSQEMNREVVFCMSFVQGEERD-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               43;
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           Youn
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           н. ү.,
                                                                                                                           Immunopathol. 4
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                                                                                                                                                           Nakamura
                                                                                                                                                                                                             Chordata; Craniata; Vertebrata; Euteleostomi; Perissodactyla; Equidae; Equus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        16.5%;
31.4%;
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                                                                                                                           e interleukin-1
48:221-231(199
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score
Pred.
                                                                                                                                                           Ν.,
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A -> G (IN REF. 2).
9D1EF8F575070586
                                                                                  Trotter G.W.,
           Watari
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Inflammatory
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mismatches
                                                                                                                                                           Nishimura Y.,
                                                                                                                           -231(1995)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         135.5;
No. 8.
          Τ.,
                                                                                                                                                                                                                                                                   update)
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                                                              cDNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        .5e-07
          Goitsuka
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                                                                                                                                    alpha
                                                                       equine
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                                                                                 Nyborg J.K.;
                                                              sequences
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                                                                                                                                                         Watari T.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CRC64;
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                                                                       interleukin-1
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                                                                                                                                                           Goitsuka R.,
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RESULT 12
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IL1B_CAPHI
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PRINTS; PR00264; INTERLEUKIN1
PROSITE; PS00253; INTERLEUKIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              InterPro; IPR000975; -.
InterPro; IPR002348; -.
Pfam; PF00340; interleukin-1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        EMBL; D42147; BAA07718.1; -. EMBL; U92481; AAC39256.1; -. EMBL; D42165; BAA22528.1; -.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               entities requires a license agreement (See http://www.isb-sib.ch/announce, or send an email to license@isb-sib.ch).
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Gene 177:11-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           "Identification of an alternatively spliced transcript of equine interleukin-1 beta.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                               PROPEP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Alternative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DOMAIN: THE SIMILARITY AMONG THE IL-1 PRECURSORS SUGGESTS THAT THE AMINO ENDS OF THESE PROTEINS SERVE SOME AS YET UNDEFINED FUNCTION.

MISCELLANGOUS: THE LACK OF A SPECIFIC HYDROPHOBIC SEGMENT IN THE PRECURSOR SEQUENCE SUGGESTS THAT IL-1 IS RELEASED BY DAMAGED CELLS OR IS SECRETED BY A MECHANISM DIFFERING FROM THAT USED FOR OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SIMILARITY: BELONGS TO THE IL-1 FAMILY.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SUBUNIT: MONOMER (BY SIMILARITY).
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                                                                                                 FLCTVPEADQPVRLTQLPENGGWNAPITDFYFQ 152
                                                                                                                                                                                      LVLSGA-----CELQAVHLNGENTNQQVVFCMSFVQGEE-----
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SWISS-PROT entry is copyright. It is produced through a collaboration een the Swiss Institute of Bioinformatics and the EMBL outstation -
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                                                                                                                                                         VQGGSQCLSCGVGQ-EPTLTLEPVNIMELYLGAKESKSFTFYRRDMGLTSSFESAAYPGW 119
                                                                                                                              LKEKNLYLSCGMKDGKPTLQLETVD-PNTYPKRKMEKRFVFNKMEIKGNVEFESAMYPNW
                                                                                                                                                                                                                                              46;
                                                                                                                                                                                                                                                          Similarity
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30.1%;
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H -> Q (IN REF. 2).
AM -> VV (IN REF. 2).
V -> M (IN REF. 2).
V -> M (IN REF. 2).
EG -> DD (IN REF. 2).
M -> V (IN REF. 2).
S -> K (IN REF. 2).
S -> K (IN REF. 2).
                                                                                                                                                                                                                                           Score 132; DB 1;
Pred. No. 2e-06;
1; Mismatches 64
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D -> N (IN REF. 2
H -> Q (IN REF. 2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Eukaryota; Metazoa;
Mammalia; Eutheria;
Bovidae; Caprinae; (
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 15-DEC-1998 (Rel. 37, Created)
15-DEC-1998 (Rel. 37, Last sequence update)
30-MAY-2000 (Rel. 39, Last annotation updat
INTERLEUKIN-1 BETA PRECURSOR (IL-1 BETA)
                                                                                                                                                                                                                                                                                                                                                                                                                                      entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    the European Bioinformatics Institute. The use by non-profit institutions as long modified and this statement is not removed.
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                                                                                                                                                                                                                                                                                    PROPEP
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PROSITE; PS00253; INTERLEUKIN_1;
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Molecular cloning
 249
                               135
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DOMAIN: THE SIMILARITY AMONG THE IL-1 PRECURSORS SUGGESTS THAT THE AMINO ENDS OF THESE PROTEINS SERVE SOME AS YET UNDEFINED FUNCTION.

MISCELLANEOUS: THE LACK OF A SPECIFIC HYDROPHOBIC SEGMENT IN THE PRECURSOR SEQUENCE SUGGESTS THAT IL-1 IS RELEASED BY DAMAGED CELLS OR IS SECRETED BY A MECHANISM DIFFERING FROM THAT USED FOR OTHER SECRETORY PROTEINS.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FUNCTION: PRODUCED BY ACTIVATED MACROPHAGES, IL-1 STIMULATES THYMOCYTE PROLIFERATION BY INDUCING IL-2 RELEASE, B-CELL MATURATION & FROLIFERATION, & FIRSTOBLAST GROWTH FACTOR ACTIVITY. IL-1 PROTEINS ARE INVOLVED IN THE INFLAMMATORY RESPONSE, BEING IDENTIFIED AS ENDOGENOUS PYROGENS, AND ARE REPORTED TO STIMULATE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SIMILARITY: BELONGS TO THE IL-1 FAMILY.
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                             QLPENGGWNAPITDF
                                                           TPTLQLEEVD-PKVYPKRNMEKRFVFYKTEIKNTVEFESVLYPNWYISTSQIEEKPVFLG
                                                                                                                                                 LKVLYLHNNQLLAGGLHAGKVIKGEEISVVPNRWLDASLSPVILGVQGGSQCLS-CGVGQ 74
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SWISS-PROT entry is copyright. It is produced through a collaboration en the Swiss Institute of Bioinformatics and the EMBL outstation -
HF -- RGGQD -- ITDF
                                                                                        EPTLTLEPVNIMELYLGAKESKSFTFYRRDMGLTSSFESAAYPGWFLCTVPEADQPVRLT 134
                                                                                                                     LKALHLISQEMSREVVFCMSFVQGEERD-----NKIPVALGIRDKNLYLSWVKKGD 189
                                                                                                                                                                              l Similarity
42; Conser
                                                                                                                                                                                                                                                                                                                                             PR00262; IL1HBGF.
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                                                                                                                                                                                                                                                                                                                                                                                              IPR000975;
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                                                                                                                                                                                                                                                        266
                                                                                                                                                                                                                                                                       114
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                                                                                                                                                                                Conservative
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31.1%;
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59F7B39BD1D4DDA5
                                                                                                                                                                                              Score 129.5; DB : Pred. No. 3.5e-06;
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RESULT 13 IL1B_PIG

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01-AUG-1992
01-AUG-1992
30-MAY-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      the European Bioinformatics Institute. There are no resuse by non-profit institutions as long as its content modified and this statement is not removed. Usage by a
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-1- FUNCTION: PRODUCED BY ACTIVATED MACROPHAGES, IL-1 STIMULATES
-1- FUNCTION: PRODUCED BY ACTIVATED MACROPHAGES, B-CELL
THYMOCYTE PROLIFERATION BY INDUCING IL-2 RELEASE, B-CELL
MATURATION & PROLIFERATION, & FIBROBLAST GROWTH FACTOR ACTIVITY.

IL-1 PROTEINS ARE INVOLVED IN THE INFLAMMATORY RESPONSE, BEING
IDENTIFIED AS ENDOGENOUS PYROGENS, AND ARE REPORTED TO STIMULATE
THE RELEASE OF PROSTAGLANDIN AND COLLAGENASE FROM SYNOVIAL CELLS.
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Gene 190 nor normal sequencing and regulation of an mRNA encoding normal sequence normal 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Cytokine; Macrophage; Mitogen; Inflammatory response; PROPER 1 114 BY SIMILARITY.
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PRINTS; PR00264; INTERLEUKIN1.
PROSITE; PS00253; INTERLEUKIN_1; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             EMBL; M86725; AAA02584.1;
PIR; JN0724; JN0724.
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Mammalia;
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DOMAIN: THE SIMILARITY AMONG THE IL-1 PRECURSORS SUGGESTS THAT THE AMINO ENDS OF THESE PROTEINS SERVE SOME AS YET UNDEFINED FUNCTION.

MISCELLANEOUS: THE LACK OF A SPECIFIC HYDROPHOBIC SEGMENT IN THE PRECURSOR SEQUENCE SUGGESTS THAT IL-1 IS RELEASED BY DAMAGED CELLS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SIMILARITY: BELONGS TO THE IL-1 FAMILY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             OR IS SECRETED BY A MECHANISM DIFFERING FROM THAT USED FOR OTHER SECRETORY PROTEINS.
                                                                                                                                                                                                                                                                                                                                                                 MVLSGALCFRMKDSALKVLYLHNNQLLAGGLHAGKVIKGEEISVVPNRWLDASLSPVILG 60
YISTSQAEQKPVFL----GNSKGRQDITDF
                                                                                                                                                                                                                                                                                                  LVLAG-----PHMLKALHLLTGDLKREVVFCMSFVQGDDSN-----NKIPYTLG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SWISS-PROT entry is copyright. It is produced through a collaboration -
                                                                      FLCTVPEADQPVRLTQLPENGGWNAPITDF 149
                                                                                                                                                 IKGKNLYLSCVMKDNTPTLQLEDID-PKRYPKRDMEKRFVFYKTEIKNRVEFESALYPNW
                                                                                                                                                                                        PF00340; interleukin-1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  P01584; 1HIB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                115
267
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (Rel. 23, Created)
(Rel. 23, Last sequence up)
(Rel. 39, Last annotation)
                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      STANDARD;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                30404 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           15.7%;
28.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                            23;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                INTERLEUKIN-1 BETA.
; 7F6B92B784D5086F CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 129;
Pred. No. '
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PRT;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    update)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DΒ
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Best Local
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SEQUENCE
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- I- SUBUNT: MONMER.
- I- DOMAIN: THE SIMILARITY AMONG THE IL-1 PRECURSORS SUGGESTS THAT THE AMINO ENDS OF THESE PROTEINS SERVE SOME AS YET UNDEFINED FUNCTION.
- I- MISCELLANEOUS: THE LACK OF A SPECIFIC HYDROPHOBIC SEGERNT IN THE PRECURSOR SEQUENCE SUGGESTS THAT IL-1 IS RELEASED BY DAMAGED CELLS
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                                                                                                                                                                                                                                                                                                                                                                                                                   use by non-profit institutions as long modified and this statement is not removed. entities requires a license agreement (See
                                                                                                                                                                                                                                                                                                                                                                                                                                                           This SWISS-PROT entry is copyright. It is produced through between the Swiss Institute of Bioinformatics and the Entropean Bioinformatics Institute. There are no resti
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             -i- FUNCTION: PRODUCED BY ACTIVATED MACROPHAGES, IL-1 STIMULATES
THYMOCYTE PROLIFERATION BY INDUCING IL-2 RELEASE, B-CELL
MATURATION & PROLIFERATION, & FIBROBLAST GROWTH FACTOR ACTIVITY.
IL-1 PROTEINS ARE INVOLVED IN THE INFLAMMATORY RESPONSE, BEING
IDENTIFIED AS ENDOGENOUS PYROGENS, AND ARE REPORTED TO STIMULATE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      01-FEB-1996 (Rel. 33, Created)
01-FEB-1996 (Rel. 33, Last sequence update)
15-DEC-1996 (Rel. 37, Last annotation updat
INTERLEUKIN-1 BETA PRECURSOR (IL-1 BETA).
                                                                                                                                                                                                                                                                       PRINTS; PR00262; IL1HBGF.
PRINTS; PR00264; INTERLEUKIN1
PROSITE; PS00253; INTERLEUKIN
                                                                                                                                                                                                                                                                                                                                                            EMBL; U19845; AAA86709.1; HSSP; P01584; 1HIB.
                                                                                                                                                                                                                                                                                                                   Pfam; PF00340;
                                                                                                                                                                                                                                                                                                                                                                                                     or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SECRETORY PROTEINS.
-!- SIMILARITY: BELONGS TO THE IL-1 FAMILY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      J. Immunol. 155:3946-3954(1995)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  nonhuman primates."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MEDLINE=96003435; PubMed=7561102;
Villinger F.J., Brar S.S., Mayne A.E.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TISSUE=Blood
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Cercopithecinae; Macaca
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                                                                                                                                                                                              PROPEP
                                                                                                                                                                                                                                                           Cytokine;
                                                                                                                                                                                                                                                                                                                                 InterPro;
                                                                                                                                                                                                                                                                                                                                              InterPro;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                "Comparative sequence analysis of cytokine genes from
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NCBI_TaxID=9544;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Macaca mulatta (Rhesus macaque).
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  120
                             178
                                                                                    134
                                                        61
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                                                                                                   1 MYLSGALCFRMKDSALKYLYLHNNQLLAGGLHAGKVIKGEEISVYPNRWLDASLSPYILG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   THE RELEASE OF PROSTAGLANDIN AND COLLAGENASE
FLCTVPEADQPVRLTQLPENGGWNA--PITDFYFQ
                            LKAKNLYLSCVLKDDKPTLQLESVD-PKNYPKKKMEKRFVFNKIEINNKLEFESAQFPNW
                                                       VQGGSQCLSCGV-GQEPTLTLEPVNIMELYLGAKESKSFTFYRRDMGLTSSFESAAYPGW 119
                                                                                                                                                                                                                                              Macrophage; Mitogen;
                                                                                                                                                                                                                                                                                                                               IPR000975; -. IPR002348; -.
                                                                                                                                                                                                                 117
269
                                                                                                                                           Conservative
                                                                                                                                                                                                                                                                                                                  interleukin-1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STANDARD;
                                                                                                                                                                                                                                                                       INTERLEUKIN_1; 1
                                                                                   ELKALHLQGQDLEQQVVFSMSFVQGEESN
                                                                                                                                                                                                                                269
                                                                                                                                                                                                                                                                                                                                                                                                                   license agreement (See http://www.isb-sib.ch/announce/
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Chordata;
Primates;
                                                                                                                                                                                                                 30481 MW;
                                                                                                                                                      15.48; 29.78;
                                                                                                                                           22;
                                                                                                                                                       Score
Pred.
                                                                                                                                                                                                                INTERLEUKIN-1 BETA,
A7CD59EBAC120EC7
                                                                                                                                                                                                                                              BY SIMILARITY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Craniata; Vertebrata; Euteleostomi; Catarrhini; Cercopithecidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PRT;
                                                                                                                                                                                                                                                         Inflammatory response;
                                                                                                                                           Mismatches
                                                                                                                                                       127;
No. 6.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Chikkala N.,
                                                                                                                                                       DB 1;
.5e-06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               8
                                                                                                                                                                                                                                                                                                                                                                                                                                              as its content
                                                                                                                                          61;
                                                                                                                                                                                                                                                                                                                                                                                                                                Usage
                                                                                                                                                                                                                 CRC64
                                                                                                                                                                   Length 269;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SYNOVIAL CELLS
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L outstation -
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Best Local
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PRINTS; PRO0264; INTERLEUKIN1.
PRINTS; PRO0264; INTERLEUKIN_1; 1.
PROSITE; PS00253; INTERLEUKIN_1; 1.
Cytokine; Macrophage; Mitogen; Inflammatory response; Pyrogen.
Cytokine; Macrophage; Mitogen; Inflamitarity.
1 116 BY SIMILARITY.
1 268 INTERLEUKIN-1 BETA.
268 CPR7266E3E2C05B4 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          IL1B_MACFA STANDARD; FKI, 200....
P79182;
15-JUL-1998 (Rel. 36, Created)
15-JUL-1998 (Rel. 36, Last sequence update)
15-DEC-1998 (Rel. 37, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entitles requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Totsuka K., Takakura H., Hashimoto O., Tatsumi M.;
Submitted (JUL-1995) to the EMBL/GenBank/DDBJ databases.
Submitted (JUL-1995) to the EMBL/GenBank/DDBJ databases.
FUNCTION: PRODUCED BY ACTIVATED MACROPHAGES, IL-1 STINULATES
THYMOCYTE PROLIFERATION BY INDUCING IL-2 RELEASE, B-CELL
MATURATION & PROLIFERATION, & FIBROBLAST GROWTH FACTOR ACTIVITY.
IL-1 PROTEINS ARE INVOLVED IN THE INFLAMMATORY RESPONSE, BEING
IDENTIFIED AS ENDOGENOUS PYROGENS, AND ARE REPORTED TO STIMULATE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Interpro; IPR000975; ...
Interpro; IPR002348; -..
Pfam; PF00340; interleukin-1; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           EMBL; D63353; BAA09677.1; -. HSSP; P01584; 1HIB.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        +
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TISSUE-Thymus;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Cercopithecinae; Macaca.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Macaca fascicularis (Crab eating macaque) (Cynomolgus monkey).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Cercopithecidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NCBI_TaxID=9541;
120 FLCTVPEADQPVRLTQLPENGGWNA--PITDFYFQ 152 :: | | | | | | | | | | | | | | |
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        INTERLEUKIN-1 BETA PRECURSOR (IL-1 BETA).
                                                              178
                                                                                                                                                134
                                                                                   61 VQGGSQCLSCGV-GQEPTLTLEPVNIMELYLGAKESKSFTFYRRDMGLTSSFESAAYPGW 119 :: : | | | | | | | | | | | | | |
                                                                                                                                                                                                                               Local Similarity
nes 46; Conserv
                                                                                                                                                                   1 MVLSGALCFRMKDSALKVLYLHNNQLLAGGLHAGKVIKGEEISVVPNRWLDASLSPVILG 60 :1:11 | | : ::||| : | | | | | |
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SUBUNIT: MONOMER.

SUBUNIT: MONOMER.

DOMAIN: THE SIMILARITY AMONG THE IL-1 PRECURSORS SUGGESTS THAT THE AMINO ENDS OF THESE PROTEINS SERVE SOME AS YET UNDEFINED FUNCTION.

MISCELLANEOUS: THE LACK OF A SPECIFIC HYDROPHOBIC SEGMENT IN THE PRECURSOR SEQUENCE SUGGESTS THAT IL-1 IS RELEASED BY DAMAGED CELLS OR IS SECRETED BY A MECHANISM DIFFERING FROM THAT USED FOR OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SIMILARITY: BELONGS TO THE IL-1 FAMILY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SECRETORY PROTEINS.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        THE RELEASE OF PROSTAGLANDIN AND COLLAGENASE FROM SYNOVIAL CELLS
                                                            LKAKNLYLSCYLKDDKPTLQLESVD-PKNYPKKKMEKREVFNKIEINNKLEFESAQFPNW 236
                                                                                                                                              LVMSGPY-----ELKALHLQGQDLEQQVVFSMSFVQGEESN---
                                                                                                                                                                                                                               Conservative
                                                                                                                                                                                                                                                 15.3%;
                                                                                                                                                                                                                               21;
                                                                                                                                                                                                                           Score 126; DB 1;
Pred. No. 8.2e-06;
11; Mismatches 62
                                                                                                                                                                                                                               62;
                                                                                                                                                                                                                                                                    Length 268;
                                                                                                                                                                                                                               Indels
                                                                                                                                           --DKIPVALG 177
                                                                                                                                                                                                                             26;
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Db 237 YISTSQAESMPVFL-----GGTRGGQDITDFTMQ 265

Search completed: June 25, 2001, 14:07:50 Job time: 146 sec

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	9.2	9.2	9.2	9.5	9.7	9.7	9.7	9.7	9.7	9.7	9.7	9.8	9.8	9.8	13.3	13.7	13.7	13.7	13.8	13.9	14.1	14.5	14.9	15.0	15.2	15.8
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ALIGNMENTS

RP SEQUENCE FROM N.A. RP SEQUENCE FROM N.A. RA BARTON J.L., Herbst R., RT "A tissue specific inte RT IL-1 cluster lacks IL-1 RL Submitted (JAN-2000) to RN [5]		SEQUENCE FROM N.A. SEQUENCE FROM N.A. MEDLINE=99443727; Pubm Mulero J.J., Pace A.M. Drmanac R., Ford J.E.; "ILLHY1: A Novel Inter Biochem. Biophys. Res.		HY1 OR ILL1. o sapiens (Human) aryota; Metazoa; malia; Eutheria; I_TaxID=9606;	Q9UBH0; 01-MAY-2000 01-MAY-2000 01-MAR-2001 FIL1 DELTA ANTAGONIST	SUL
SEQUENCE FROM N.A. SEQUENCE FROM N.A. Barton J.L., Herbst R., Bosisio D., Nicklin M.J.H.; Pa tissue specific interleukin-1 receptor antagonist homolog from the "A tissue specific Interleukin-1 receptor antagonist homolog from the "IL-1 cluster lacks IL-1, IL-1Ba and IL-1Bra activities."; Submitted (JAN-2000) to the EMBL/GenBank/DDBJ databases.	e F.S., Symons J.A., Nicklin M.J.H.; cerleukin-1 receptor antagonist homolog from the II-1ra, IL-18 and IL-18ra activities."; o the EMBL/GenBank/DDBJ databases.	<pre>led=10512743; , Nelken S.T., Loeb D.B., Correa T.R., leukin-1 Receptor Antagonist Gene."; Commun. 263:702-706(1999).</pre>	<pre>led=10625660; R., Ketchem R.R., Kubin M., Garka K.E., nd the IL-1 Superfamily."; i9-1175(2000).</pre>	Chordata; Craniata; Vertebrata; Euteleostomi; Primates; Catarrhini; Hominidae; Homo.	(TremBLrel. 13, Created) (TremBLrel. 13, Last sequence update) (TremBLrel. 16, Last annotation update) (INTERLEUKIN-1 LIKE PROTEIN.1) (INTERLEUKIN-1 RECEPTOR HOMOLOG 1) (INTERLEUKIN-1 DELTA).	(Y; PRT; 155 AA.

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RESULT
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OPPOSE
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EMBL; AF366094; AAF02757.1; --
EMBL; AJ242737; CAB59822.1; --
EMBL; AJ242738; CAB59823.1; --
EMBL; AJ242738; CAB59823.1; --
EMBL; AJ271338; CAB67704.1; --
EMBL; AF216693; AAF76981.1; --
EMBL; AF230377; AAF91274.1; --
HSSP; P18510; IILR,
HSSP; P18510; IILR
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Q9QYY1;
01-MAY-2000
                                                                                        MEDLINE=20209405; PubMed=10744718;
Kumar S., McDonnell P.C., Lehr R., Tierney
Griswold D.E., Capper E.A., Tal-Singer R.,
                                                                                                                                                                                                              SEQUENCE FROM N.A.

Barton J.L., Nicklin M.J.H.;

Barton J.L., Nicklin M.J.H.;

"IL-IL1: A Novel Member of the Interleukin-1 Gene Family

in Trophoblasts and Macrophages.";

Submitted (OCT-1999) to the EMBL/GenBank/DDBJ databases.
                                                                                                                                                                                                                                                                                                                                                                                                                                               01-MAY-2000 (TrEMBLrel. 13, Created)
01-MAY-2000 (TrEMBLrel. 13, Last sequence update)
01-MAR-2001 (TrEMBLrel. 16, Last annotation update)
IL-1L1 PROTEIN (INTERLEUKIN-1 HOMOLOG 3).
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                                                                                                                                                                                                                                                                                                                                                                                                          Eukaryota; Metazoa;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Receptor.
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PROSITE; PS00253; INTERLEUKIN_1;
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Kastelein R.A.;
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                                                                                                                                                                                                                                                                                                                                                       Mammalia; Eutheria;
NCBI_TaxID=10090;
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[6]
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                        Identification and initial characterization the interleukin-1 family.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              121 LCTVPEADQPVRLTQLPENGGWNAPITDFYFQQCD 155
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           121
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  Biol.
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Motazoa; Chordata;
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       Chem.
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  1-1 family.";
275:10308-10314(2000)
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01-OCT-2000 (TrEMBLrel. 1
01-MAR-2001 (TrEMBLrel. 1
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Kastelein R.A.;
"Novel IL-1 ligands IL-1d ar
Submitted (FEB-2000) to the
EMBL; AF230378; AAF91275.1;
                                                                                                                                                                                                                                                                                                                                                                                                       Interpro; IPR000975; -.
pfam; pF00340; ILI; I.
PRINTS; PR00264; INTERLEUKIN1.
PROSITE; PS00253; INTERLEUKIN_1;
SMART; SM00125; ILI; 1.
SEQUENCE 156 AA; 17136 MW; A4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mus musculus (Mouse).
Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Rodentia;
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EMBL; AF200495; AAF69251.1;
HSSP; P18510; IILR:
InterPro; IPR000975; ...
Pfam; PF00340; IL1; 1.
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PROSITE; PS00253; INTERLEUKIN_1;
SMART; SM00125; III; 1
SEQUENCE 155 AA; 17004 MW; A/
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE FROM N.A.
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                      LCTVPEADQPVRLTQLPENGGWNAPITDFYFQQCD
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LCTSPEADQPVRLTQIPEDPAWDAPITDFYFQQCD
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                                                                                                                                                                                                                                                                                                                                                                                                       17136 MW; A4D1EE2F93CF77A7 CRC64;
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Pred. No. 3.8e-67;
4; Mismatches 10
                                                                                                                                                                                                                                                                                       Score 734; DB 11;
Pred. No. 3.8e-67;
4; Mismatches 10;
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Sciurognathi; Muridae;
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Best Local S
Matches 73
MEDLINE=97146044; PubMed=8992991; Jenkins J.K., Drong R.F
                                                                                                                                                                                                                                                                                                                                                                                                                                            Q14628
Q14628;
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NON_TER
SEQUENCE
                                                                                                                                                                                                                                                                         Homo sapiens (Human).
Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Primates;
                                                                                                                                                                                                                                                                                                                                                            01-NOV-1996 (TrEMBLrel 01, 01-NOV-1996 (TrEMBLrel 01, 01-MAR-2001 (TrEMBLrel 16, INTRACELLULAR IL-1 RECEPTOR
                                                                                                                                                                                            SEQUENCE FROM N.A. MEDLINE=95355865; PubMed=7629520;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE FROM N.A. MEDLINE=98183404; PubMed=9514884;
                                                                                           "Cloning and characterization receptor antagonist.";
J. Exp. Med. 182:623-628(1995)
                                                                                                                                                      Muzio M., Polentarutti N., Sironi
Introna M., Mantovani A., Colotta
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Pfam; PF00340; IL1; 1.
PRINTS; PR00264; INTERLEUKIN1.
PROSITE; PS00253; INTERLEUKIN_1; 1.
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             01-MAY-2000
01-MAR-2001
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01-MAY-2000
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HSSP; P18510; 1ITN
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73; Conserv
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143 AA;
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Pred. No. 5.6e
13; Mismatches
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ANTAGONIST TYPE II.
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RECEPTOR ANTAGONIST VARIANT (FRAGMENT).
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                                                                                                                                                                                                                                                                           Craniata; Vertebrata; Euteleostomi; Catarrhini; Hominidae; Homo.
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Best Local
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MEDLINE-98209757; PubMed-9550387;

Gabay C., Porter B., Fantuzzi G., Arend W.

"Mouse IL-1 receptor antagonist isoforms:
and protein expression of intracellular is
distribution of secreted and intracellular
                                                                                                                                                                                                                                                                                                     INTRACELLULAR (Mouse).
Mus musculus (Mouse).
Larvota; Metazoa; Chordata; (
                                                                                                                                                                                                                                                                                                                                                                070207 PRELIMINARY;
070207;
01-AUG-1998 (TrEMBLrel. 07,
01-AUG-1998 (TrEMBLrel. 07,
01-MAR-2001 (TrEMBLrel. 16,
                                                              PROSITE; PS00253; INTERLEUKIN_1; SMART; SM00125; IL1; 1. SEQUENCE 159 AA; 17995 MW; A
                                                                                                  Pfam; PF00340; IL1; 1.
PRINTS; PR00264; INTERLEUKIN1.
ProDom; PD002536; -; 1.
                                                                                                                                                  J. Immunol. 159:5905-5913(1997).
EMBL; AF001795; AAC15251.1; -.
HSSP; P18510; 1IRA.
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PRINTS; PR00254; INTERLEUKIN1.
ProDom; PD002536; -; 1.
PROSITE; PS00253; INTERLEUKIN_1; 1.
SMART; SM00125; ILI; 1.
SEQUENCE 180 AA; 19897 MW; 624A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Submitted (JUL-1996) to the EMBL; X84348; CAA59087.1; -. EMBL; U65590; AAB92269.1; -. HSSP; P18510; IIRP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Arend W.P., Smith M.F. Jr.;
"Intracellular IL-1 receptor antagonist and inducible regulatory regions.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE FROM N.A. Slightom J.L.;
                                                                                                                                                                                          vivo."
                                                                                                                                                                                                                                                                              SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                    NCBI_TaxID=10090;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           InterPro; IPR000975; -.
                                                                                                                                       InterPro; IPR000975; -.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               J. Immunol.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  154
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          126
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                EADQPVRLTQLPENGGWNAPITDFYFQQ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FRMKDSALKVLYLHNNQLLAGGLHAGKVIKGEEISVVPNRWLDASLSP--VILGVQGGSQ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CLSCVKSGDETRLQLEAVNITDLSENRKQDKRFAFIRSDSGPTTSFESAACPGWFLCTAM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CLSC-GVGQEPTLTLEPVNIMELYLGAKESKSFTFYRRDMGLTSSFESAAYPGWFLCTVP 125
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FRIWDVNQKTFYLRNNQLVAGYLQGPNVNLEEKIDVVP-----IEPHALFLGIHGGKM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  EADQPVSLTNMPDEG - - - VMVTKFYFQE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              38.8%;
             37.5%;
                                                                                                                                                                                                   intracellular isoform and and intracellular IL-1 rece
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     13;
                                                                                                                                                                                                                                                                                                                                                      Last sequence update)
Last annotation updat
RECEPTOR ANTAGONIST.
                                                                                                                                                                                                                                                                                                                                                                                         Created)
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             Score
Pred.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 319.5;
Pred. No. 7.
                                                                                                                                                                                                                                                                                                              Craniata; Vertebrata; Euteleostomi; Sciurognathi; Muridae; Murinae; Mus
                                                                                                                                                                                                                                                                                                                                                                                                                      PRT;
                                                              A7CD35D137846222 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  624A1574C2334229 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           153
                                                                                                                                                                                                                                        Arend W.P.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  178
             308.5; DB 1
No. 8.4e-24;
                                                                                                                                                                                                                                                                                                                                                                                                                      159
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      promoter: cell type-specific
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                , DB 4
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     49;
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Similarity

Conservative

15;

Mismatches

49;

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13;

Gaps

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DB 11;

Length

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OPERING
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Best Local
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                                                                                                                                                                                                                                                                                                                                                             Q9GKK2 PRELIMINARY; PRT; 176 AA.
Q9GKK2;
01-MAR-2001 (TrEMBLrel. 16, Created)
01-MAR-2001 (TrEMBLrel. 16, Last sequence update)
01-MAR-2001 (TrEMBLrel. 16, Last annotation update)
INTERLEUKIN-1 RECEPTOR ANTAGONIST.
                                                                                                                                                                                                   Canis familiaris (Dog).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata;
Mammalia; Eutheria; Carnivora; Fissipedia; Canidae;
NCBI_TaxID=9615;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Receptor.
SEQUENCE
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SEQUENCE FROM N.A.

Campbell S.E., Nasir L., Argyle D.J., Gault E., B
"Canine II-1 Receptor Antagonist cDNA Sequence.";
Submitted (DEC-1999) to the EMBL/GenBank/DDBJ dat
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Tursiops truncatus (Atlantic bottle-nosed dolphin).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Cetartiodactyla; Cetacea; Odontoceti; Delphin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Inoue Y., Itou T., Sakai T.;
"Cloning and Sequencing of a Bottle-Nosed Dolphin Interleukin-1 Receptor Antagonist.";
Submitted (FEB-2000) to the EMBL/GenBank/DDBJ databases.
EMBL, AB038268; BAB11806.1;
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SEQUENCE FROM N.A.
Thou T.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NCBI_TaxID=9739;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CLSC-GVGQEPTLTLEPVNIMELYLGAKESKSFTFYRRDMGLTSSFESAAYPGWFLCTVP 125
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DQPVRLTQLPENGGWNAP - - ITDFYFQQ
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DRPVSLTNTPEE----PLIVTKFYFQE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FRIWDTNQKTFYLRNNQLIAGYLQGPNIKLEEKIDMVP----IDLHSVFLGIHGGKLCL
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FRIWDVNQKTFYLRNNQLVAGYLQGPNTKLEEKIDVVP-----IEPHAMFLGIHGGKL 90
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            177 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        19923 MW;
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48.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        6FD19A06C09B131B
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                 databases
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Best Local S
Matches 68
                                                                                                                                                                                                                                       Matches
                                                                                                                                                                                                                                                              Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Q9UHA5; PRELIMINARY;
Q9UHA5;
01-MAY-2000 (TrEMBLrel. 1
01-MAY-2000 (TrEMBLrel. 1
01-MAR-2001 (TrEMBLrel. 1
077771;
077771;
01-NOV-1998
                                                                                                                                                                                                                                                                                                                                                                                          Sims J.E.;
"Four New Members Expand the TL-1 Superfamily.";
J. Biol. Chem. 275:1169-1175(2000).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          EMBL; AF216526; AAG36777.1; Receptor. SEQUENCE 176 AA; 19938 MY
                                                                                                                                                                                                                                                                                                  Interpro; IPR000975; -.

Pfam; PF00340; IL1; 1.

PRINTS; PR00264; INTERLEUKIN1.

PROSITE; PS00253; INTERLEUKIN_1; UNKNOWN_1.

SMART; SM00125; IL1; 1.

SEQUENCE 157 AA; 17702 MW; 7A54F3D7557A
                                                                                                                                                                                                                                                                                                                                                                         J. Biol. Chem. 275:1169-117/
EMBL; AF201833; AAF25213.1;
HSSP; P10749; 2MIB.
                                                                                                                                                                                                                                                                                                                                                                                                                                     MEDLINE=20092888; PubMed=10625660; Smith D.E., Renshaw B.R., Ketchem
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Eutele
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE FROM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NCBI_TaxID=9606;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens (Human)
                                                                                      128
                                                                                                             122
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                                                                                                                                      83
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                                                                                                                                                                                                                                                  Local Similarity
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                                                                                                            CTVPEADQPVRLTQLPENGGWNAPITDFYFQQCD 155
                                                                                                                                     GKDLCLFCAEIQGKPTLQLKEKNIMDLYVEKKAQKPFLFFHNKEGSTSVFQSVSYPGWFI 127
                                                                                                                                                GGSQCLSCG-VGQEPTLTLEPVNIMELYLGAKESKSFTFYRRDMGLTSSFESAAYPGWFL 121
                                                                                                                                                                                      YAIRDSRQMVWVLSGNSLIAAPL--SRSIKPVTLHLIACR--DTEFSDKEKGNMVYLGIK 67
                                                                                                                                                                                                             FRMKDSALKVLYLHNNQLLAGGLHAGKVIKGEEISVVPNRWLDASLSP-----VILGVQ 62
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      EADQPVRLTQLPENGGWNAPITDFYFQQ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CLACVKSGDETRLQLEAVNITDLSKNKDQDKRFTFILSDSGPTTSFESAACPGWFLCTAL 150
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CLSC-GVGQEPTLTLEPVNIMELYLGAKESKSFTFYRRDMGLISSFESAAYPGWFLCTVP 125
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FRMKDSALKVLYLHNNQLLAGGLHAGKVIKGEEISVVPNRWLDASLSP--VILGVQGGSQ 66
                                                                                     ATSTTSGQPIFLTK - - ERGITNN - - TNFYLDSVE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              EADRLVSLTNRPEEA---MMVTKFYFQK 175
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FRIWDVNQKTFYLRNNQLVAGYLQGSNTKLEEKLDVVP-----VEPHAVFLGIHGGKL 90
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                68;
                                                                                                                                                                                                                                       Conservative
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                                                                                                                                                                                                                                                                                                 17702 MW; 7A54F3D7557A3EE3 CRC64;
                                                                                                                                                                                                                                                 24.4%;
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16,
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Last sequence update)
Last annotation updat
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Pred. No. 5.3e
13; Mismatches
                                                                                                                                                                                                                                      Score 200.5; DB 4
Pred. No. 8.2e-13;
4; Mismatches 61
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                                                                                                                                                                                                                                                                                                                                                                                                                                      R.R.,
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                                                                                      157
                                                                                                                                                                                                                                                                                                                                                                                                                                      Kubin M.,
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                                                                                                                                                                                                                                      61;
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Q9JLA2
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     01-OCT-2000
01-OCT-2000
01-MAR-2001
                                                       Debets R., Timans J., Zurawski S., Bazan J.F., Kastelein R.A "Novel IL-1 family member IL-Le responds through the orphan related protein 2; response is antagonized by IL-1d."; Submitted (NOV-1999) to the EMBL/GenBank/DDBJ databases. EMBL; AF200493; AAF69249.1; -. EMBL; AF200697; AAG35671.1; -. EMBL; AF206697; AAG35671.1; -. ITTELETO: IPRO00975; -.
                                                                                                                                                                                                                                                                                                                                           SEQUENCE FROM N.A.

MEDLINE=20209405; PubMed=10744718;
Kumar S., McDonnell P.C., Lehr R., Tier
Griswold D.E., Capper E.A., Tal-Singer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NON_TER
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Antagonist (IL-Ira) Gene.";
Submitted (JUN-1998) to the EMBL/GenBank/DDBJ databases.
EMBL; AF072476; AAC62237.1; -.
                  PRINTS;
                                         InterPro; IPR000975; -
Pfam; PF00340; IL1; 1.
                                                                                                                                                                                                                                                                                        the
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PROSITE; PS00253; INTERLEUKIN_1;
SMART; SM00125; IL1; 1.

NON_TER 1 1
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Eukaryota; Metazoa; Cho
Mammalia; Eutheria; Per
NCBI_TaxID=9796;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mus musculus (Mouse)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             01-OCT-2000 (TrembLrel.
01-OCT-2000 (TrembLrel.
01-MAR-2001 (TrembLrel.
INTERLEUKIN-1 HOMOLOG 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Q9JLA2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Q9JLA2
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Dhar A.K., Thompson M.S.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               01-NOV-1998
01-MAR-2001
                                                                                                                                                                                                                     SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                        "Identification and initial characterization
                                                                                                                                                                                                                                                                                                                                    Young P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NCBI_TaxID=10090;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE FROM N.A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   141
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         11
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      81 EPVNIMELYLGAKESKSFTFYRRDMGLTSSFESAAYPGWFLCTVPEADQPVRLTQLPENG 140
                                                                                                                                                                                                                                                                   Biol.
                                                                                                                                                                                                                                                                                     interleukin-1
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  SM00125;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       39;
                  PR00264;
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                                                                                                                                                                                                                                                                   Chem.
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(TrEMBLrel.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRELIMINARY;
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                                                                                                                                                                                                                                                              n-1 family.";
275:10308-10314(2000)
                    INTERLEUKIN1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  8215
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Chordata;
Rodentia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Chordata; Craniata; Vert
Perissodactyla; Equidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              53
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15, Last sequence update)
16, Last annotation update)
(INTERLEUKIN-1 EPSILON).
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16, Last annotation update)
ANTAGONIST SECRETORY FORM (F
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Pred. No. 4.2e-1
0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Craniata; Vertebrata; Euteleostomi; Sciurognathi; Muridae; Murinae; Mus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRT;
                                                                                                                                                                                                                                                                                                                                                  , Tierney
inger R.,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Vertebrata;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Alcivar-Warren
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Ā
                                                                                                                                                                                                                                                                                                                                                  L., Tzimas !
Wells G.I.,
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                                                                                                                                                                                                                                                                                                        novel members
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                                                                                                                                                                                                                                                                                                                                                  Doyle M.L.,
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RESULT 13
Q9UHA6
ID Q9UHA6
AC Q9UHA6
DT 01-MAY
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Q9UHA7
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local
  FILI ZETA.
FILI ZETA.
Homo saplens (Human).
Eukaryota; Metazoa; Cheria; Portaria; Port
                                                                                                                                                                                                                                                                                                                                                                                                                                                            Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Matches
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Q9UHA6;
Q1-MAY-2000
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Q9UHA7;
01-MAY-2000
01-MAY-2000
01-MAR-2001
                                                                                                   01-MAY-2000
01-MAR-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PRINTS; PR00264; INTERLEUKIN1.
SMART; SM00125; IL1; 1.
SEQUENCE 158 AA; 17684 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       J. Biol. Chem. 275:1169-1175(2000).
EMBL; AF201831; AAF25211.1; -.
HSSP; P18510; 11RA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE FROM N.A.
MEDLINE=20092888; PubMed=10625660;
Smith D.E., Renshaw B.R., Ketchem
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              InterPro; IPR000975; .
Pfam; PF00340; IL1; 1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Eukaryota;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NCBI_TaxID=9606;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    "Four New Members Expand the IL-1 Superfamily.";
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                                                                                                                                                                                                                                                                                                                    114 AAYPGWFLCTVPEADQPVRLTQLPENGGWNAPITDF 149
                                                                                                                                                                                                                                                                                             121
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           12
                                                                                                                                                                                                                                                                                                                                                                                                                   SPVILGVQGGSQCLSCG-VGQEPTLTLEPVNIMELYLGAKESKSFTFYRRDMGLTSSFES 113
                                                                                                                                                                                                                                                                                                                                                                          NPIYLGLNGLNLCLMCAKVGDQPTLQLKEKDIMDLYNQPEPVKSFLFYHSQSGRNSTFES 120
                                                                                                                                                                                                                                                                                             VAFPGWFIAVSSEGGCPLILTQ--ELG--KANTTDF 152
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        IAVCSKGSCPLILTQ--ELG--EIFITDF
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Similarity
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                                                                                                   (TrEMBLrel. 13, (TrEMBLrel. 13, (TrEMBLrel. 16,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 160 AA;
                                                                                                                                                                                       PRELIMINARY;
                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
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Primates;
                       Primates;
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                                         Chordata;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                22.5%;
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36.2%;
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13,
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                                                                                                     Last
Last
                                                                                                                                              Created)
                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 185.5;
Pred. No. 2.8e
13; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Created)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 186.5;
Pred. No. 2.2
                     Craniata; V
Catarrhini;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Craniata; Vo
Catarrhini;
                                                                                                                                                                                       PRT;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  469AC84306B0E280 CRC64;
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                                                                                                                                                                                       192
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                                           Vertebrata; Euteleostomi;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Vertebrata; 1 i; Hominidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Kubin
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.2e-11;
                                                                                                   on update)
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                       Hominidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DB 4;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                            35;
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QPNZH6
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Best Local S
Matches 51
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                O9NZH6 PRELIMINARY; PRT; 218 AA.
O9NZH6;
O9NZH6;
O1-OCT-2000 (TrEMBLrel. 15, Created)
O1-OCT-2000 (TrEMBLrel. 15, Last sequence update)
O1-MAR-2001 (TrEMBLrel. 16, Last annotation update)
INTERLEUKIN-1 HOMOLOG 4 (IL-1x PROTEIN) (INTERLEUKIN-1-RELATED PROTEIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            "Four New Members Expand the IL-1 Superfamily.";
J. Biol. Chem. 275:1169-1175(2000).
EMBL; AF201832; AAF25212.1;
HSSP; P18510; 1ITN.
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SEQUENCE FROM N.A.
MEDLINE-20092888; PubMed-10625660;
Smith D.E., Renshaw B.R., Ketchem
                                                                                                                                                                                                                                              Pan G., Risser P., Mao W., Baldwin D.T., Zhong A.W., Yansura Lewis L., Elgenbrot C., Henzle W.J., Vandlen R., Filvaroff E. "IL-1H, an interleukin-1-related protein that binds IL-18 rec
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TISSUE-FETAL LUNG, FETAL TESTIS, FETAL B-CITISSUE-FETAL LUNG, FETAL TESTIS, FETAL B-CITISSUE-FETAL B-CITIS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   InterPro; IPR000975; -.
Pfam; PF00340; IL1; 1.
PRINTS; PR00264; INTERLEUKIN1
SMART; SM00125; IL1; 1.
                                                                  EMBL; AF200496; AAF69252.1;
EMBL; AF167368; AAG29344.1;
EMBL; AF251118; AAG14420.1;
                                                                                                                                                                                                                                                                                                                                                                                                                                             Manoj P.P., Mantovani A., Muzio M.;
Submitted (JUL-1999) to the EMBL/GenBank/DDBJ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                the
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Homo sapiens (Human).
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                                                                                                                                                                                   Submitted (APR-2000) to the
                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE FROM N.A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE FROM N.A.
                                                                                                                                                                                                                          lRrp."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 "Identification and initial characterization the interleukin-1 family.";
J. Biol. Chem. 275:10308-10314(2000).
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PF00340;
                               IPR000975; -.
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36.7%;
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Pred. No. 7.4e-10;
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Wells G.I., Doy
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O9HBF3;
O1-MAR-2001 (TrEMBLrel. 16, Created)
O1-MAR-2001 (TrEMBLrel. 16, Last sequence update)
O1-MAR-2001 (TrEMBLrel. 16, Last annotation update)
INTERLEUKIN-1-RELATED PROTEIN LONG ISOFORM.
                                                                                                                                                                                                                                                                                                                                                                                     Homo sapiens (Human).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                          Submitted (APR-2000) to the EMBL/GenBank/DDBJ databases EMBL; AF251119; AAG14421.1; . SEQUENCE 218 AA; 24138 MW; 76E09C35093DEA63 CRC64;
                                                                                                                                                                                                                                                                                                  SEQUENCE FROM N.A.

Pan G., Risser P., Mao W., Baldwin D.T., Zhong A.W., Yansura
Pan G., Risser P., Mao W., Baldwin D.T., Zhong A.W., Yansura
Lewis L., Eigenbrot C., Henzle W.J., Vandlen R., Filvaroff E.
"IL-1H, an interleukin-1-related protein that binds IL-18 rec
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PRINTS; PR00264; INTERLEUKIN1.
SEQUENCE 218 AA; 24126 MW;
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                                                                                                                                        FRMKDSALKVLYLHNNQLLAGGLHAGKVIKGEEISVVPNRWLDASL---SPVILGVQGGS
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ICTSCNCNEPVGVTDKFEN
                           LCTVPEADQPVRLTQLPEN
                                                       FCLYCDKDKGQSHPSLQLKKEKLMKL-AAQKESARRPF1FYRAQVGSWNMLESAAHPGWF
                                                                                  QCLSC--GVGQ-EPTLTLEPVNIMELYLGAKES--KSFTFYRRDMGLTSSFESAAYPGWF 120
                                                                                                                FSIHDQDHKVLVLDSGNLIA--VPDKNYIRPEIFFALASSLSSASAEKGSPILLGVSKGE
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Pred. No. 8.7e
21; Mismatches
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